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Perfect score:
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Aae21296
Adi16993
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_	Ad196466	Abw00803	Adc86445	Abp59266	Abp96696	Adi16997	Adi16636	Adi16991	Aae21288	Abp95617	Aae17074	Aa:104371	Aae12794	Aag64294	Adc17728	Aay87664	Adc12766	Aay30162	Adw02566	Aay30161
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## ALIGNMENTS

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WPI; 199
N-PSDB;
                                                                                                                                                                                                                                                  AAY30159 standard; protein; 322
                                                                                                        Ahmad
                                                                                                                                              16-DEC-1998;
                                                                                                                                                         01-JUL-1999
                                                                                                                                                                                                           Human dorsal root receptor 1 hDRR1.
                                                                                                                                                                                                                      20-MAR-2003
26-NOV-1999
                                                                                                                                                                                                                                        AAY30159;
                                                                                                                   (ASTR )
                                                                                                                                                                     W09932519-A1
                                                                                                                                                                               Homo sapiens
                                                                                                                                    22-DEC-1997;
                                                                                      1999-405162/34.
DB; AAZ10067.
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                                                                                                                                                                                          root receptor; dorsal root ganglia; G-protein coupled receptor;
central nervous system; CNS; anaesthesia; analgesia; neuron; pain.
                                                                                                                   ASTRA
ASTRA
                                                                                                       Banville
                                                                                                                                                                                                                      (revised)
(first entry)
                                                                                                                   PHARMA INC.
AB.
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This is the human dorsal root receptor 1 (hDRR1) protein sequence. It is a G protein coupled receptor that is expressed preferentially in dorsal root ganglia. hDRR1 can be used to create antibodies against hDRR1. The dorsal root ganglia area of the central nervous system (c is densely innervated with primary or afferent neurons involved in transmission, modulation and sensation of pain. The DRR's which are expressed in this region of the CNS may be used for assays for the identification of new agents for anaesthesia and analgesia. (Updated 20-MAR-2003 to correct PA field.)

(Updated

g

(CNS)

This

Claim 6;

Page 39-41; 72pp; English.

and human dorsal root receptors and related polynucleotides, useful identifying agents for anesthesia and analgesia.

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ADW46672
ADW46672
ADW46672
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AC ADW4
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Best Local
                                                                                                                                                WPI; 200
N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sensory neuron-specific G protein-coupled receptor 1; SNSR1; g-protein coupled receptor; gpc; diagnosis; cardiovascular disease; gastrointestinal disease; liver disease; cancer; neoplasm; inflammation; hematological disease; respiratory disease; neurological disease; genitourinary disease; cardiovascular-gen.; cardiant; vasotropic; antiarrhythmic; antiarreriosclerotic; hypotensive; gastrointestinal-gen.; antiinflammatory; antiulcer; hepatotropic; antiallergic; dermatological; thyromametor; antipressive; antianemic; cytostatic; hemostatic; antiasthmatic; respiratory-gen.; cns-gen.; antiparkinsonian; nootropic; antiasthmatic; cerebroprotective; nephrotropic; uropathic; receptor.
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                        Screening for therapeutic agents, useful for treating e.g., cardiovascular disorders, comprises contacting a test compound with sensory neuron-specific G protein-coupled receptor 1 (SNSR1) polypes
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                                                                                                                                                                                                                                                                                                                                                      16-JUN-2003; 2003EP-00013598
                                                                                                                                                                                                                                                                                                                                                                                                               04-JUN-2004; 2004WO-EP006078
   detecting binding
                                                                                                                                                2005-057894/06.
DB; ADW46671.
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Pred. No. 2.4e-175;
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Disclosure; SEQ ID NO 2; 122pp; English

The invention relates to a method of screening for therapeutic agents C creating diseases associated with sensory neuron-specific G protein-C coupled receptor (SNSR1). The method comprises contacting a test compound with SNSR1 polypeptide or polynucleotide and detecting the compound with SNSR1 polypeptide or polynucleotide and detecting the concentration, (b) in the absence of the test compound, or (c) at a concentration of the test compound. Also described are (i) a different concentration of the test compound. Also described are (i) a method of diagnosing a disease defined above in amammal, (ii) a method of diagnosing a disease defined above in comprising the SNSR1 polypeptide. (iii) a SNSR1 polypeptide or polynucleotide, or a cherry polynucleotide of the test compound. Also described are (i) a method of diagnosing the SNSR1 polypeptide activity such as a small molecule, an ANA molecule, an antisense oligonucleotide, a polypeptide, an antibody, or a ribozyme, and (iv) a method for the preparation of a pharmaceutical composition or treating the above diseases. The SNSR1 regulatory compounds are useful for preparing a pharmaceutical composition for treating diseases such as cardiovascular disorders, gastroinestinal and liver diseases, cancer disorders, inflammatory diseases, hematological disorders, respiratory diseases, hematological disorders, respiratory diseases, meurological disorders, or unological disorders, respiratory diseases include fast the regulation of SNSR1 cut diseases and chronic pancreatitis. Inflammatory diseases include atherosclerosis in the systemic lupus expthematory diseases include atherosclerosis, stroke, and Althoric pancreatitis. Inflammatory diseases include atopic diseases, and chronic pancreatitis. Inflammatory diseases include atopic diseases, service diseases incl

Sequence 322 AA;

Length

밁 ঠ 밁 ঠ 밁 δ 밁 S 밁 Query Match Best Local S Matches 322 181 181 121 121 13 5 مبر 1 MDPTIPVLGTKLTPINGREETPCYNQTLSFTGLTCIISLVALTGNAVVLWLLGCRMRRNA Similarity CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSDFIT VSIYILNLVAANFLFLSGHIIFSPLPLINIRHPISKILSPVMTFPYFIGLSMLSAISTER VSIYILNLVAANFLFLSGHIIFSBLPLINIRHPISKILSPVMTFPYFIGLSMLSAISTER MDPTIPVLGTKLTPINGREETPCYNQTLSFTGLTCIISLVALTGNAVVLWLLGCRMRRNA VDEGGGWLPQETLELSGSKLEQ RIHLDWKVLFCHVHLYSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPE CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSDFIT RIHLDWKVLFCHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPE IAWLVFLCVVLCGSSLVLLVRILCGSRKWPLTRLYVTILLTVLVFLLCGLPFGIQWALFS IAWLVFLCVVLCGSSLVLLVRILCGSRKMPLTRLYVTILLTVLVFLLCGLPFGIQWALFS Conservative 100.0%; Score 1691; DB 9; 100.0%; Pred. No. 2.4e-175; tive 0; Mismatches 0; Indels ç Gaps 120 120 60 60 300 240 180 180

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301

VDEGGGWLPQETLELSGSKLEQ

322

300 300

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RESULT 3
AAY90762
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Best Local Similarity 97.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence represents a human-derived G protein-coupled protein designated hHI77213V, which is isolated from the human hippocampus. The G protein-coupled receptor can be used for preventing, treating and diagnosing genetic diseases associated with G protein-coupled protein, and for regulating cellular functions. The protein can be used to prevent and treat disorders associated with G protein-coupled protein gration. It can also be used to identify G protein-coupled protein ligands and generating antibodies and antisera against the protein. It is also useful in constructing recombinant receptor protein expression systems, developing receptor-binding assay systems and screening drug candidates, and can be used as a probe in the genetic diagnosis of G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human-derived G protein-coupled protein and encoding nucleic acid, useful e.g. in determining ligands and treatment of diseases associated with dysfunction of the protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 322 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           candidates, and can be used as a protein-coupled protein disorders
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IAMLVFLCVVLCGSSLVLLVRILCGSRKMPLTRLYVTILLTVLVFLLCGLPFGIQMALFS
                                                                                                                    CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSDFIT
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                                                                                CLSILWDIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSDFIT
                                                                                                                                                                                                                                                                                                                    MDSTIPVLGTELTPINGREETPCYKQTLSFTGLTCIVSLVALTGNAVVLWLLGCRMRRNA
                                                                                                                                                                                                        VSIYILNLVAADFLFLSGHIICSPLRLINIRHPISKILSPVMTFPYFIGLSMLSAISTER
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Pred. No. 1.5e
4; Mismatches
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L.5e-170;
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27-FBB-2001;
27-FBB-2001;
27-FBB-2001;
02-MAR-2001;
14-MAR-2001;
14-MAR-2001;
14-MAR-2001;
14-MAR-2001;
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08-FEB-2001;
09-FEB-2001;
15-FEB-2001;
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31-JAN-2001;
31-JAN-2001;
31-JAN-2001;
                                                                                                                                              16-MAR-2001;
16-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human; NOVX; cardiomyopathy; atherosclerosis; cancer; diabetes; inflammation; autoimmune disorder; allergy; blood disorder; acquired immunodeficiency syndrome; AIDS; obesity; asthma; immunoglobulin (Ig)A nephropathy; cirrhosis; arthritis; Alzheimer's disease; infection; str.
                                                                                                                                                                    15-MAR-2001;
15-MAR-2001;
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2001US-0276448P.
2001US-0276450P.
2001US-027639P.
2001US-0276768P.
2001US-0278775P.
2001US-0278775P.
2001US-0278778984P.
2001US-028014P.
2001US-028014P.
2001US-0280149.
2001US-0285133P.
2001US-0285133P.
2001US-0285137P.
2001US-028537P.
2001US-028837P.
2001US-028837P.
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2001US-0268974P.
2001US-0271665P.
2001US-0271839P.
2001US-0271855P.
2001US-02727889.
2001US-0273046P.
2001US-0275925P.
2001US-0275947P.
2001US-0275949P.
2001US-0275949P.
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; 2001US-0266975P.
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2001US-0267459P.
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180

180

300

240 240

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Cd diagnostic and prognostic assays and furthermore in the treatment of cdiverse pathological conditions. The present invention describes novel thuman and murine NOVX proteins, as well as methods to modulate their cd expression using antisense oligos, ribozymes and peptide nucleic acids. The NOVX polypeptides, polynucleotides and antibodies are useful in treating or preventing NOVX-associated disorders, e.g. cardiomyopathy, catherosclerosis, cancer and diabetes. Furthermore, they may be used in ctreating or preventing diseases such as inflammation, autoimmune cd disorders, allergies, blood disorders, acquired immunodeficiency syndrome cd disorders, acquired immunodeficiency syndrome cd antipates, blood disorders, acquired immunodeficiency syndrome cd include policy, accordingly, these molecules have many activities including crystatic, anti-HIV, antidiabetic, antiarteriosclerotic, antialergic, antisathmatic, nephrotropic, antibaterial, virucide, antiparasitic, continuous protestive, antiallergic, antisathmatic, hepartotropic, antibaterial, virucide, antiparasitic, relaxant and anticonvulsant. In addition, they are useful in screening cassays to identify small molecules that modulate or inhibit, for example, continuous as in chromosome mapping, tissue typing, preventive medicine and confirmation. This polypeptide is a homologue of a human NOVX protein confirmation.
                                                                                                                                                                                                               Best Loc
Matches
                                                                                                                                                                                                                                                                 Query Match
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18-JUN-2001;
19-JUN-2001;
13-AUG-2001;
16-AUG-2001;
16-AUG-2001;
21-AUG-2001;
21-AUG-2001;
31-AUG-2001;
31-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention relates to a novel nucleic acids, and encoded polypeptides thereof, which have properties related to the stimulation of biochemical or physiological responses in a cell, tissue, organ or organism. Specifically, it refers to the use of biologically active fragments for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-SEP-2001;
12-SEP-2001;
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atherosclerosis, or
pharmacogenomics.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New NOVX polypeptides and nucleic acids, useful for preventing or treating NOVX-associated disorders, e.g. cancer, cardiomyopathy, atherosclerosis, or diabetes, and in chromosome mapping, tissue typing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 530; 1498pp; English.
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Gangolli EA, Padigaru M, Ander
K, Taupier NJ, Gusev VY, Cc
K, Grosse WM, Alsobrook JP, I
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                           VSIYILNLVAANFLFLSGHIIFSPLFLINIRHPISKILSFVMTFFYFIGLSMLSAISTER
  VSIYILNLVAADFLFLSGHIICSPLRLINIRHPISKILSPVMTFPYFIGLSMLSAISTER
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2001US-0312020P.
2001US-0312889P.
2001US-0313390P.
2001US-0315470P.
2001US-0316447P.
2001US-0318115P.
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2001US-03181290P.
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2001US-03181290P.
2001US-0323379P.
2001US-0323379P.
2001US-0333308P.
2001US-0333701P.
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ilarity 97.5%;
Conservative
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Pred. No. 1.5e-170;
4; Mismatches 4;
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Anderson DW, Rastelli L, Miller CE;
VY, Colman SD, Wolenc AR, Pena CEA;
JP, Lepley DM, Rieger DK, Burgess CE;
                                                                                                                                                                                                               4;
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RESULT 5
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05-FEB-2001

07-FEB-2001

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08-FEB-2001

15-FEB-2001

26-FEB-2001

27-FEB-2001

27-FEB-2001

27-FEB-2001

14-MAR-2001

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15-MAR-2001

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-JAN-2002; 2002WO-US002785.
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2001US-0266767P
2001US-0266767P
2001US-026795P
2001US-0267859P
2001US-0268974P
2001US-0271859P
2001US-0271859P
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2001US-0273046P
2001US-0275947P
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2001US-0278779P
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2001US-0266406P.
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CC diagnostic and prognostic assays and furthermore in the treatment of CC diverse pathological conditions. The present invention describes novel CC expression using antisense oligos, ribozymes and peptide nucleic acids. The NOVX polypeptides, polymucleotides and antibodies are useful in CC treating or preventing NOVX-associated disorders, e.g. cardiomyopathy, CC atherosclerosis, cancer and diabetes. Furthermore, they may be used in CC treating or preventing diseases such as inflammation, autoimmune CC disorders, allergies, blood disorders, acquired immunodeficiency syndrome CC disorders, Alzheimer's disease, infections, stroke, muscular dystrophy and epilepsy. Accordingly, these molecules have many activities including CC cytostatic, cardiant, antinflammatory, immunosuppressive, antialtergic, CC haemostatic, anti-HIV, antidiabetic, antiarteriosclerotic, anorectic, antiasthmatic, nephrotropic, antiastratic, they are useful in screening CC assays to identify small molecules that modulate or inhibit, for example, ceff the invention. This polypeptide is a homologue of a human NOVX protein CC fe the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-MAY-2001;
29-MAY-2001;
30-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-JUN-2001
19-JUN-2001
13-AUG-2001
16-AUG-2001
16-AUG-2001
21-AUG-2001
28-AUG-2001
28-AUG-2001
07-SEP-2001
07-SEP-2001
12-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                This invention relates to a novel nucleic acids, and encoded polypeptides thereof, which have properties related to the stimulation of biochemical or physiological responses in a cell, tissue, organ or organism. Specifically, it refers to the use of biologically active fragments for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New NOVX polypeptides and nucleic acids, useful for preventing or treating NOVX-associated disorders, e.g. cancer, cardiomyopathy, atherosclerosis, or diabetes, and in chromosome mapping, tissue typing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 535; 1498pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        pharmacogenomics.
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Gangolli EA, Padigaru M, Anderson DW, I
h VL, Taupier RJ, Gusev VY, Colman SD,
K, Grosse WM, Alsobrook JP, Lepley DM,
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2001US-0294047P

2001US-029404P

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2001US-0298959P

2001US-0312020P

2001US-0312020P

2001US-0312089P

2001US-0312908P

2001US-0313390P

2001US-0315470P

2001US-0318118P

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2001US-031814P

2001US-031814P
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son DW, Rastelli L, Miller CE;
lnan SD, Wolenc AR, Pena CEA;
epley DM, Rieger DK, Burgess C
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Burgess CE;

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Best Local Sir Matches 314; Query Match Best Local Similarity

Conservative

97.4**%**; 97.5**%**;

Score 1647; Pred. No. 1 Mismatches

.5e-170; DB 5;

Length 322;

0

Gaps

0

Query Match

97.4%;

Score 1647;

멂 7;

Length 322

322

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ARESULT 6
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                                                            The invention relates to a novel polynucleotide encoding a guanosine triphosphate-binding protein coupled receptor (GPCR). A polynucleotide of the invention may have a use in gene therapy. The polynucleotide and polypeptide are useful for preparing a composition for treating a patient in need of increased or suppressed activity or expression of the guanosine triphosphate-binding protein coupled receptor. The protein sequences shown in ADC85549-ADC87617 represent GPCR's of the invention.
   Sequence 322 AA;
                                                                                                                                                                                                                                                                                                                              Claim
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N-PSDB; ADC86820.
                                                                                                                                                                                                                                                                                                                                                                                                                         New polynucleotide, useful for preparing a composition for treating a patient in need of increased or suppressed activity or expression of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Suwa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY. (ADSC-) CENT ADVANCED SCI & TECHNOLOGY INCUBATI
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                                                                                                                                                                                                                                                                                                                           SEQ ID NO 1274; 28pp; English.
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The invention relates to a non-human mammal that carries a DNA integrated with a foreign HI7T213 or its mutant gene, or a part of it. The non-human animal is particularly a rat. Such gene shows phenotypes of e.g. cataract onset, transient skin rash and proliferation-promoting activity. The foreign HI7T213 gene is a gene that encodes a G protein-coupled receptor (GPCR) protein HI7T213. The protein, its encoded DNA and constructed transgenic animals are useful for clarifying pathological mechanism,
                                                                                                                                                                                                                                                            HI7T213 protein, encode pathological mechanism, preventives or remedies
                                                                                                                                                                                           Claim 3; SEQ ID NO 3; 161pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                N-PSDB; ADO44605
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human HI7T213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADO44604 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (TAKE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transgenic; G protein-coupled receptor; GPCR; ophthalmological;
ic; nephrotropic; antiinflammatory; dermatological; analgesic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSDFIT
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                                                                                                                                                                                                                                                            encoded DNA and transgenic animals for clarifying hanism, developing therapeutic methods and screening emedies for related diseases e.g. cataract, cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein;
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4; Mismatches 4;
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Best Local Simi
Matches 314;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            developing therapeutic methods and screening preventives or related diseases e.g. cataract, cancer, and dermatitis. The sequence represents a human HI7T213 protein.
                                              Human-derived G protein-coupled protein and encoding nucleic acid, us e.g. in determining ligands and treatment of diseases associated with dysfunction of the protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 322 AA;
                                                                                                                                                                                                                                              genetic
                                                                                                                                                                                                                                                         Human;
                                                                                                                                                                                                                                                                         Human G protein-coupled receptor hHI7T213
                                                                                                                                                                                                                                                                                                                               AAY90761 standard;
                                                                                    N-PSDB;
                                                                                             WPI;
                                                                                                               Watanabe
                                                                                                                                                    01-OCT-1998;
                                                                                                                                                                      30-SEP-1999;
                                                                                                                                                                                        13-APR-2000
                                                                                                                                                                                                          WO200020455-A1
                                                                                                                                                                                                                             Homo
                                                                                                                                                                                                                                                                                            18-AUG-2000
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                                                                                             2000-303747/26
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                                                                                                                                                                                                                                              G protein-coupled receptor; hippocampus; c disease; cellular function regulation.
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                                                                                    AAA29811.
                                                                                                                                  TAKEDA
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No. 1.5e-170;
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                                                                                                                                                                                                                                                       diagnosis;
                                                                                                                                                                                                                                                                           NO:1.
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The present sequence represents a human-derived designated hH17T213, which is isolated from the

G protein-coupled human hippocampus.

protein. The G

Claim 1;

Page 90-91; 97pp;

Japanese.

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Best Local Sim
Matches 313;
          27-NOV-2000;
12-DEC-2000;
20-FEB-2001;
20-FEB-2001;
06-APR-2001;
06-APR-2001;
06-APR-2001;
06-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein-coupled receptor can be used for preventing, treating and diagnosing genetic diseases associated with G protein-coupled protein, and for regulating cellular functions. The protein can be used to preve and treat disorders associated with G protein-coupled protein gene dysfunction. It can also be used to identify G protein-coupled protein ligands and generating antibodies and antisera against the protein. It also useful in constructing recombinant receptor protein expression systems, developing receptor-binding assay systems and screening drug candidates, and can be used as a probe in the genetic diagnosis of G protein-coupled protein disorders
                                                                                                                                                                                                                                    Human; G-protein coupled receptor; GPCR; hRUP28; hRUP29; hRUP30; hRUP31; hRUP32; hRUP33; hRUP35; hRUP36; hRUP37.
                                                                                                                                                                                                                                                                        Human G protein coupled receptor hRUP37.
                                                                                                                                                                                                                                                                                                   11-OCT-2002
                                                                                                                                                                                                                                                                                                                                                   ABJ04077 standard;
                                                                                                                                     26-NOV-2001; 2001WO-US044386
                                                                                                                                                                                      WO200242461-A2
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2000US-025366P.
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2001US-0282335P.
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97.2%;
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Pred. No. 5.2e-170;
5; Mismatches 4;
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Best Local S
Matches 313
                                                                                                                            Human; mas-related gene; G-protein coupled receptor; receptor; sensory perception; pain; analgesic; MrgX3
 04-MAY-2000; 2000US-0202027P
                        04-MAY-2001; 2001WO-US014519.
                                                                                                                                                                  Human MrgX3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention provides the protein and coding sequences of several human G-protein coupled receptors (GPCRs). These can be use the identification of candidate compounds as receptor agonists or agonists having applicability as therapeutic agents. The present is a GPCR protein of the invention
                                                                                                                                                                                             01-JUL-2002
                                                                                                                                                                                                                      AAE21296;
                                                                                                                                                                                                                                               AAE21296 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel endogenous and non-endogenous versions of G protein-coupled receptor useful for identification of candidate compounds as receptors agonists or antagonists for use as therapeutic agents.
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313; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61 VSIYILNLVAANFLFLSGHIIFSPLPLINIRHPISKILSPVMTFPYFIGLSMLSAISTER
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97.2%;
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Pred. No. 5.2e
5; Mismatches
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5.2e-170;
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03-NOV-2000;
19-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to Mrg (mas-related gene) protein, which is a G-protein coupled receptor and drg-12 protein, which is a receptor. The invention is useful for identifying compounds that bind to it, especially agonists or antagonists. Administration of an agent (e.g. the identified agonist) that increases the expression of Mrg in a mammal may be used for treating impaired sensory perception in a mammal, especially pain. The antagonist may also be useful for treating impaired sensory perception in a mammal. The present sequence is human MrgX3 protein
                       human; NOVX; cardiomyopathy; atherosclerosis; cancer; diabetes; inflammation; autoimmune disorder; allergy; blood disorder; acquired immunodeficiency syndrome; AIDS; obesity; asthma; immunoglobulin (Ig)A nephropathy; cirrhosis; arthritis; Alzheimer's disease; infection; str.
                                                                                                                                                                  Human
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2001US-0285493P.
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5.2e-170;
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                                                             ev VT, Spytek KA, Zerhusen BI
Gangolli EA, Padigaru M, And
h VL, Taupier RJ, Gusev VY,
K, Grosse WM, Alsobrook JP,
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2001US-029324P.
2001US-0312020P.
2001US-0312889P.
2001US-03133908P.
2001US-0315470P.
2001US-0316447P.
2001US-0318115P.
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2001US-0265412P.
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2001US-0267459P.
2001US-0267753P.
2001US-0267753P.
2001US-0271649P.
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2001US-02721839P.
2001US-0275947P.
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2001US-0288337P.
2001US-02988504P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2002WO-US002785
                                                                                                                   Zerhusen BD,
                                                                                                                   Patturajan M,
                                                                  jan M, Shimkets RA;
Rastelli L, Miller CE;
Wolenc AR, Pena CEA;
I, Rieger DK, Burgess C
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NOVX polypeptides

and

nucleic

acids,

useful

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preventing

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Anderson DW, F VY, Colman SD, C JP, Lepley DM,

GB;

2002-706998/76.

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RESULT 12
ADI16998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention relates to a novel nucleic acids, and encoded polypeptides CC thereof, which have properties related to the stimulation of biochemical cor physiological responses in a cell, tissue, organ or organism. CC Specifically, it refers to the use of biologically active fragments for cdiverse pathological conditions. The present invention describes novel thuman and murine NOVX proteins, as well as methods to modulate their corporating NOVX proteins, as well as methods to modulate their corporating NOVX-associated disorders, e.g. cardiomyopathy, corporating novy-associated disorders, e.g. cardiomyopathy, corporating or preventing Novy-associated disorders, e.g. cardiomyopathy, corporated in a lergies, blood disorders, acquired immunodeficiency syndrome (AIDS), obesity, asthma, immunoglobulin (Ig)A nephropathy, cirrhosis, cardiars, allergies, blood disorders, acquired immunodeficiency syndrome (AIDS), obesity, asthma, immunoglobulin (Ig)A nephropathy, cirrhosis, cardiart, antiinflammatory, immunosuppressive, antialergic, and policy and apilepsy. Accordingly, these molecules have many activities including cytostatic, anti-HIV, antidiabetic, antiarteriosclerotic, anorectic, neuroprotective, nootropic, antiarthritic, hepatotropic, antiallergic, corporated in danticonvulsant. In addition, they are useful in screening corporated in chromosome mapping, tissue typing, preventive medicine and cophramacogenomics. This polypeptide is a homologue of a human NOVX protein coffice of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Human NOVX protein homologue SeqID 534.
                                                                                                  ADI16998 standard;
                                     15-APR-2004
                                                                     ADI16998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; SEQ ID NO 529; 1498pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            pharmacogenomics.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        treating NOVX-associated disorders, e.g. cancer, cardiomyopathy, atherosclerosis, or diabetes, and in chromosome mapping, tissue typing or
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                                                                                                                                                                                                                                                                                                                                                                                                                                            RIHLDWKVLFCHVHLVSIFLSALMSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPE
                                                                                                                                                                                                                                                                                                                                  IAWLVFLCVVLCGSSLVLLVRILCGSRKMPLTRLYVTILLTVLVFLLCGLPFGIQWALFS
                                                                                                                                                                                                                                                                                                                                                                                                CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSDFIT
                                                                                                                                                                                                 VDEGGGWLPQETLELSGSKLEQ 322
                                                                                                                                                                                                                                               RIHLDWKVLFCHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPE
                                                                                                                                                                                                                                                                                                              IAWLVFLCVVLCGSSLVLLVRILCGSRXMPUTRLYVTILLTVLVFLLCGLPFGIQWALFS
                                                                                                                                                                                                                                                                                                                                                                             CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGADSVWCETSDFIT
                                                                                                                                                                                VDEGGGWLPQETLELSGSRLEQ 322
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                                 (first entry)
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                                                                                                  protein; 322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97.18;
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Pred. No. 5
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5.2e-170;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 322;
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15-MAR-2001;
16-MAR-2001;
16-MAR-2001;
20-MAR-2001;
26-MAR-2001;
29-MAR-2001;
29-MAR-2001;
30-MAR-2001;
11-APR-2001;
11-APR-2001;
11-APR-2001;
20-APR-2001;
20-MAY-2001;
23-MAY-2001;
23-MAY-2001;
23-MAY-2001;
23-MAY-2001;
13-MAY-2001;
11-APR-2001;
23-MAY-2001;
23-MAY-2001;
21-JUN-2001;
11-JUN-2001;
11-JUN-2001;
11-JUN-2001;
11-JUN-2001;
11-JUN-2001;
11-JUN-2001;
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08-FEB-2001;
09-FEB-2001;
15-FEB-2001;
26-FEB-2001;
27-FEB-2001;
27-FEB-2001;
02-MAR-2001;
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31-JAN-2001;
31-JAN-2001;
31-JAN-2001;
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                       (CURA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-JAN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Alzheimer's disease; infection; str.
                       CURAGEN CORP
                                          2001US-0267823P

2001US-0268974P

2001US-0271834P

2001US-0271834P

2001US-0271834P

2001US-02727846P

2001US-0275947P

2001US-0275947P

2001US-0276450P

2001US-0276450P

2001US-0276450P

2001US-0276450P

2001US-02767676P

2001US-02767676P

2001US-0278652P

2001US-0278652P

2001US-0278775P

2001US-0278678P

2001US-0278652P

2001US-0278652P

2001US-0278652P

2001US-0278652P

2001US-0278652P

2001US-0279884P

2001US-0289947P

2001US-0289479P

2001US-0299477P

2001US-0299477P

2001US-0312808P

2001US-0312808P

2001US-03112908P

2001US-03113390P

2001US-03118115P

2001US-0318115P

2001US-0318115P

2001US-0318115P

2001US-0318115P

2001US-0318115P

2001US-0318115P

2001US-0318115P

2001US-031874P

2001US-031874P
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2001US-0267459P.
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Spytek KA,

Zerhusen BD,

Patturajan M,

Shimkets RA;

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New NOVX polypeptides and nucleic acids, useful for preventing or treating NOVX-associated disorders, e.g. cancer, cardiomyopathy, atherosclerosis, or diabetes, and in chromosome mapping, tissue typing
                                                                                                                                                                                                                                      pharmacogenomics.
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                                                                                                                                                                                                                                                                                                  Gerlach
                                                                                                                                                                                                                                                                                   n VL, Taur K, Grosse WM,
                                                                                                                                                                                                                        SEQ ID
                                                                                                                                                                                                                                                                                         lli EA, Padigaru M, Anderson DW,
Taupier RJ, Gusev VY, Colman SD,
rosse WM, Alsobrook JP, Lepley DM,
                                                                                                                                                                                                                       NO 534; 1498pp; English.
                                                                                                                                                                                                                                                                                          Rastelli L,
Wolenc AR,
1, Rieger DK,
                                                                                                                                                                                                                                                                                          Pena CEA;
Burgess CE;
                                                                                                                                                                                                                                                                                                         Miller CE;
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This invention relates to a novel nucleic acids, and encoded polypeptides thereof, which have properties related to the stimulation of biochemical CC thereof, which have properties related to the stimulation of biochemical CC specifically, it refers to the use of biologically active fragments for CC diagnostic and prognostic assays and furthermore in the treatment of CC diverse pathological conditions. The present invention describes movel thuman and murine NOVX protesins, as well as methods to modulate their CC expression using antisense oligos, ribozymes and peptide nucleic acids. CC The NOVX polypeptides, polynuclectides and antibodies are useful in CC treating or preventing NOVX-associated disorders, e.g. cardiomyopathy, CC atherosclerosis, cancer and diabetes. Furthermore, they may be used in CC treating or preventing diseases such as inflammation, autoimmune CC disorders, allergies, blood disorders, acquired immunodeficiency syndrome CC (AIDS), obesity, asthma, immunoglobulin (Ig)A nephropathy, cirrhosis, CC arthritis, Alzheimer's disease, infections, stroke, muscular dystrophy CC and epilepsy. Accordingly, these molecules have many activities including CC cytostatic, anti-HIV, antidiabetic, antiarteriosclerocic, anorectic, neuroprotective, nootropic, antiartritic, hepatotropic, antiallergic, relaxant and anticonvulsant. In addition, they are useful in screening CC relaxant and anticonvalsant. In addition, they are useful in screening cused as in chromosome mapping, tissue typing, preventive medicine and of the invention.

CC of the invention. pharmacogenomics. This of the invention.

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Matches
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         121
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                                                                                                                                                                              Similarity
                       CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMPCDFLFSGANSVWCETSDFIT
                                                                           VSIYILNLVAANFLFLSGHIIFSPLFLINIRHPISKILSPVMTFPYFIGLSMLSAISTER
          CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGADSVWCETSDFIT
                                                              VSIYILNLVAADFLFLSGHIICSPLRLINIRHPISKILSPVMTFPYFIGLSMLSAISTER
                                                                                                              MDSTIPVLGTELTPINGREETPCYKQTLSFTGLTCIVSLVALTGNAVVLWLLGCRMRRNA
                                                                                                                             MDPTIPVLGTKLTPINGREETPCYNQTLSFTGLTCIISLVALTGNAVVLWLLGCRMRRNA
                                                                                                                                                                 Conservative
                                                                                                                                                                             97.1%;
97.2%;
                                                                                                                                                                 Score 1642; DB 5;
Pred. No. 5.2e-170;
5; Mismatches 4;
                                                                                                                                                                                          Length 322;
                                                                                                                                                                   ٥,
                                                                                                                                                                     Gaps
                                       180
                                                                120
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             180
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문 S 241 VDEGGGWLPQETLELSGSKLEQ 322 RIHLDWKVLFCHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPE RIHLDWKVLFCHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPE 밁

181 181

300

240

240

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RESULT 13 ABP81750

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Human G protein-coupled receptor MrgX3 protein SEQ ID NO:674
                                                                                                      04-MAR-2003
                                                                                                             ABP81750 standard; protein;
                                                                                                       (first entry)
                                                                                                             322
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og P

g protein-coupled receptor; GPCR; antigenic peptide; gene therapy;
g protein-coupled receptor modulator; antibody; immune-related disease;
growth-related disease; cell regeneration-related disease; AIDS; cancer;
immunological-related cell proliferative disease; autoimmune disease;
Alzheimer's disease; atherosclerosis; infection; osteoarthritis; allergy;
osteoporosis; cardiomyopathy; inflammation, Crohn's disease; diabetes;
graft versus host disease; Parkinson's disease; multiple sclerosis; pain;
psoriasis; anxiety; depression; schizophrenia; dementia; memory loss;
mental retardation; epilepsy; asthma; tuberculosis; obesity; nausea;
hypertension; hypotension; renal disorder; rheumatoid arthritis; trauma;

Homo WO200261087-A2 sapiens.

08-AUG-2002.

19-DEC-2001; 2001WO-US050107.

19-DEC-2000; 2000US-0257144P.

LIFESPAN BIOSCIENCES INC

2003-046718/04.

GC,

Roush CL,

New isolated antigenic peptides e.g., for G protein-coupled receptors (GPCR), useful for diagnosing and designing drugs for treating conditions in which GPCRs are involved, e.g. AIDS, Alzheimer's disease, cancer or

Disclosure; Fig 1; 523pp; English.

cc any one of 1601 sequences (see ABB22019 to ABB23619) of 12-24 amino co acids. Also described: (1) an assay for the detection of a particular G CC protein-coupled receptor (GPCR) or a candidate polypeptide in a sample; cc and (2) an isolated antibody having high specificity and high affinity or CC avidity for a particular GPCR. (1) can be used as GPCR modulators and in CC gene therapy. The antigenic peptides for GPCRs are useful in detecting an CC antibody against a particular GPCR, and in the production of specific cc antibodies. The peptides and antibodies are also useful for detecting the GPCRs and antibodies are also useful for detecting the CC gresence or absence of corresponding GPCRs. The antigenic peptides for CC GPCRs and antibodies are useful for diagnosing and designing drugs for treating immune-related diseases, growth-related diseases, cell corresponding the corresponding GPCRs. The antigenic peptides for CC diseases, or autoimmune diseases, e.g. AIDS, Alzheimer's disease, cell costeoarthritis, osteoporosis, cancer, cardiomyopathy, chronic and acute CC diseases, osteoporosis, cancer, cardiomyopathy, chronic and acute CC disease, Parkinson's disease, multiple sclerosis, pain, psoriasis, cancer, depression, schizophrenia, dementia, mental retardation, memory consideration, renal disorders, rheumatoid arthritis, trauma, ulcers, or CC used in immunoassays and immunodiagnosis. ABZ42523 to ABZ42869 encode corrections of the nreacer involved. The antibodies may be corrected to the proteins given in ABP81675 to ABB82018, which are used in the corrected to the proteins given in ABP81675 to ABB82018, which are used in the GPCR proteins given in ABP81675 to ABP82 exemplification of the present invention The present invention describes antigenic peptides (I) comprising: any one of 1601 sequences (see ABP82019 to ABP83619) of 12-24 amino (a)

Sequence

Query Match

Score 1642; В Length 322;

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RESULT 14
ADH08535
ID ADH08
XX ADH08
XX ADH08
XX ADH08
XX MAS-r
XX AD18c
XX MAS-r
XX Disc
X
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The present invention relates to an isolated mas-related gene D (MrgD) polypeptide. The MrgD polypeptides are useful as therapeutics or for identifying compounds, i.e. agonists or antagonists, that alter pain perception in a mammal. The compounds are useful for treating impaired sensory perception, e.g. chronic intractable pain or neuropathic pain, promoting wound healing, restoring normal sensitivity following injury, or treating ocular conditions, particularly those associated with pressure such as glaucoma. The Mrg genes or proteins may be used as
                                                                                                                                                                                                                                                                            New mas-related gene D polypeptides, useful as therapeutics or in identifying agonists or antagonists that alter pain perception in a mammal for treating impaired sensory perception, e.g. chronic intractable pain or neuropathic pain.
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                                                                                                                                                                                                                                Disclosure; SEQ ID NO 31; 220pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (CALY ) CALIFORNIA INST OF TECHNOLOGY.
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perception;
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glaucoma; Mrg.
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5; Mismatches 4;
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Best Local Simi
Matches 313;
                                                                                          HI7T213 protein, encoded pathological mechanism, o preventives or remedies f
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HI7T213; transgenic; G protein-coupled receptor; GPCR; ophthalmological; cytostatic; nephrotropic; antiinflammatory; dermatological; analgesic; vulnerary; neuroprotective; human; receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           molecular probes for the detection of cells or tissues related to involved with sensory perception. The present sequence represents (Mrg subfamily) protein.
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                                                                                          d DNA and transgenic animals for clarifying developing therapeutic methods and screening for related diseases e.g. cataract, cancer, a
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Query Match 97.1%; Score 1642; DB 8; Length 322; Best Local Similarity 97.2%; Pred. No. 5.2e-170; Matches 313; Conservative 5; Mismatches 4; Indels
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RESULT 2 TVRTAS transform C;Species C;Date: 3 C;Accessi R;Young, R;Young, Proc. Nat A;Title:	RESULT 1 A39485 Cransformin Cransformin Cransformin Cransformin Cransformin Cransformin Cransformin Rimonot Ri		0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
2 es: Rat 31-Dec sion: A atl. Ac :: Chara	RESULT 1 A39485 A39485 Cransforming protein Crispecies: Homo sapie Crispecies: A39485 R;Monnot, Crispeciol. 5, 1 A;Title: Cloning and A;Reference number: A39485 A;Residues: Preliminary A;Residues: Lype: DNA A;Residues: Lype: Lype: DNA A;Residues: Lype: Lype: Lype: Lype: Lype: Lype: Lype: Lype: Lype: L		194.5 194.5 192.5 192.5 193.6 199.1 199.1 188.5 188.5 186.5 186.5 180.5 180.5
RESULT 2 TVRTAS transforming protein mas - C;Species: Rattus norvegie C;Date: 31-Dec-1989 #seque C;Accession: A31816 R;Young, D.; O'Neill, K.; Proc. Natl. Acad. Sci. U.& A;Title: Characterization	SULT 1 9485 9485 Tansforming protein (mr species: Homo sapiens Date: 28-Feb-1992 #seq Accession: A39485 Accession: A3948		11.55 11.55 11.4.55 10.77
protein mas - rat ittus norvegicus ( ic-1989 #sequence_ A31816 O'Neill, K.; Jess cad. Sci. U.S.A. acterization of t	rotein (mrg) - human to sapiens (man) -1992 #sequence_revisio 39485 Meber, V.; Stinnakre, J 1. 5, 1477-1487, 1991 mg and functional chara mber: A39485; MUID:9213 39485 iminary te: DNA 378 <mon* 26.8%;="" 378="" <mon*="" protein-coupled="" recep<="" recepto="" sco="" td=""><td></td><td>35 35 36 36 36 36 36 36 36 36 36 36 36 36 36</td></mon*>		35 35 36 36 36 36 36 36 36 36 36 36 36 36 36
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at) 31-Dec-198 31-Dec-198 Wigler, M. -5342, 1988 as oncogene	ion 17-Apr-1993 #text_change 09-Jul-20 J.; Bihoreau, C.; Teutsch, B.; Corvol accterization of a novel mas-related g 30997; pMID:1723144  UNIPARC:UPI000003B44C; GB:S78653; NID or; transmembrane protein core 453; DB 2; Length 378; ed. No. 5.9e-30; Mismatches 80; Indels 46; Gaps Mismatches 80; Indels 46; Gaps [:             : :                 : :	ALIGNMENTS	
9 #t	7-Apr-1993 #text_change 09-Jul-20 Bihoreau, C.; Teutsch, B.; Corvol rization of a novel mas-related g 7; PMID:1723144  ARC:UPI000003B44C; GB:S78653; NID transmembrane protein 453; DB 2; Length 378; No. 5-9e-30; matches 80; Indels 46; Gaps mitches 80; Indels 46; Gaps FIGLSMLSAISTERCUSILWPIWYHCRRPRYLS :		
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A;Gene: mas
C;Superfamil
C;Keywords:
F;31-47/Doma
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A;Accession: A31816
A;Molecule type: mRNA
A;Residues: 1-324 <YOU>
A;Cross-references: UNIPROT:P12526; UNIPARC:UPI0000043DE7; GB:J03823; NID:g205313; C;Genetics:
                                                                                       F;105-135/Domain: transmembrane #status predicted <TM3>
F;150-172/Domain: transmembrane #status predicted <TM4>
F;156-214/Domain: transmembrane #status predicted <TM5>
F;186-214/Domain: transmembrane #status predicted <TM6>
F;255-250/Domain: transmembrane #status predicted <TM6>
F;258-286/Domain: transmembrane #status predicted <TM7>
                                                                                                                                                                                  A;Map position: 6q24-6q27
C;Superfamily: mas transforming protein
C;Keywords: G protein-coupled receptor; glycoprotein; proto-oncogene; transforming
F;31-61/Domain: transmembrane #status predicted <TM1>
F;66-97/Domain: transmembrane #status predicted <TM2>
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C;Genetics:
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                                                                                                                                                                                                                                                                             A;Gene: GDB:MAS1
A;Cross-references: GDB:120166; OMIM:165180
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A; Residues: 1-325 < YOU>
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Matches 107
                                                                        ,225-250/Domain: transmembrane
,258-286/Domain: transmembrane
,5,16,22,272/Binding site: carb
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                 Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GSRKMPLTRLYVTILLTVLVFLLCGLPFGIQWALFSRIHLDWKVLFCHVHLVSIFLSALN
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     Conservative
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                                                                         site: carbohydrate
                    25.4%;
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   63;
   Score 429.5;
Pred. No. 4.4e
53; Mismatches
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                                       DB 1; Length 325;
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     27;
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Local

Similarity

107;

Conservative

55;

Mismatches

99;

Indels

23;

Gaps

37 ISLVALTGNAVVLWLLGCRMRRNAVSIYILNLVAANFLFLSGHIIFS-----PLPLINIR

205 161

GSRKMPLTRLYVTILLTVLVFLLCGLPFGIQMALFSRIHLDWKVLFCHVHLVSIFLSALN : | : | : | : | : | : | : | : | : |

101

HYTIVTLSVTFLFGYNTGLYLLTAISVERCLSVLYPIWYTSHRPKHQSAFVCALLCALSC 160 HPISKILSPVMTFPYFIGLSMLSAISTERCLSILWPIWYHCRRPRYLSSVMCVLLWALSL 

151

LRSILEWMFCDFLFSGANSVWCETSD-----FITI-AWLVFLCVVLCGSSLVLLVRILC LVTTMEYVMC--IDSGEES--HSRSDCRAVIIFIAILSFLVFTPLMLVSSS-ILVVKIRK

215 204 92 41

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A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-87,'I',89-324 <RES>
A;Cross-references: UNIPARC:UPI0000029856; EMBL:X67735;
C;Genetics:
A;Gene: mas
C;Superfamily: mas transforming protein
C;Keywords: G protein-coupled receptor; proto-oncogene;
                                                                                                                                                                                                                                                                               R;Metzger, R.; Bader, M.; Ludwig, T.; Berberich, C.; Bunneme FBBS Lett. 357, 27-32, 1995
A;Title: Expression of the mouse and rat mas proto-oncogene A;Reference number: I48647; MUID:95094925; PMID:8001672
A;Accession: I48647
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N;Alternate names: mas proto-oncogene protein; probable G protein-coupled receptor C;Species: Mus musculus (house mouse)
C;Date: 10-Apr-1996 #sequence revision 19-Apr-1996 #text_change 09-Jul-2004
C;Accession: S51001; 148647; S29619
R;Metzger, R.; Bader, M.; Ludwig, T.; Berberich, C.; Bunnemann, B.; Ganten, D.
FEBS Lett. 357, 27-32, 1995
A;Title: Expression of the mouse and rat mas proto-oncogene in the brain and periph A;Accession: S51001
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S51001
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R;Metzger, R.; Bader, M.; Ludwig, T.; Berberich, C.; Bunnemann, B.; Ganten,
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                       Query Match
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25.0%; Score 422.5; DB 2; 37.7%; Pred. No. 1.7e-27;
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                                                                             proto-oncogene; transmembrane
                             Length
                                                                                                                                                                                     NID: 953011;
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A35639
G protein-coupled rec
C;Species: Rattus nor
C;Date: 22-Jan-1993 #
C;Accession: A35639
R;Ross, P.C.; Figler,
                    A; Molecule type: DNA
A; Residues: 1-355 <BEC>
A; Cross-references: UNIPROT: P21109; UNIPARC: UPI000012D4ED;
A; Cross-references: UNIPROT: P21109; UNIPARC: UPI000012D4ED;
A; Lee, J; Kuang, W.J.; Rice, G.C.; Wood, W.I.
J. Immunol. 148, 1261-1264, 1992
A; Title: Characterization of complementary DNA clones encod A; Reference number: A46483; MUID: 92148149; PMID:1737938
A; Accession: A46483
A; Accession: A46483
A; Molecule type: mRNA
A; Molecule type: mRNA
A; Molecule type: mRNA
A; Residues: 1-355 <LEE>
A; Residues: 1-355 <LEE
A; Residues: 1-355
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A;Title: RTA, a candidate G protein-coupled receptor: cloning, sequencing, and tiss A; Reference number: A35639; MUID:90222168; PMID:2109324
A;Reference number: A35639
A;Residues: A35639
A;Residues: 1-343 <ROS>
A;Residues: 1-343 <ROS>
A;Cross-references: UNIPROT:P23749; UNIPARC:UPI0000046F10; GB:M35297; NID:g206809; C;Keywords: G protein-coupled receptor; glycoprotein; membrane protein C;Keywords: G protein-coupled receptor; glycoprotein; membrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Accession: JQ1231; A46483
R;Beckmann, M.P.; Munger, W.E.; Kozlosky, C.; VandenBos, T.; Price Biochem. Biophys. Res. Commun. 179, 784-789, 1991
A;Title: Molecular characterization of the interleukin-8 receptor A;Reference number: JQ1231; MUID:91378994; PMID:1898400
A;Accession: JQ1231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                interleukin-8 receptor - rabbit
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 31-Mar-1992 #sequence_revision 31-Mar-1992
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A;Cross-references: UNIPARC:UPI000012D4ED;
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Species: Rattus norvegicus (Norway rat)
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                                                                                                                                                                                           DNA clones encoding PMID:1737938
GB:M82873; NID:g165440; PIDN:AAA31376.1;
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R;Thomas, K.M.; Pyun, H.Y.; Navarro, J.
J. Biol. Chem. 265, 20061-20064, 1990
A;Title: Molecular cloning of the fMet-Leu-Phe receptor from neutrophils.
A;Reference number: A23669; MUID:91056034; PMID:1700779
A;Accession: A23669
A;Accession: A23669
A;Accession: A23669
A;Residues: 1-354 < CTHO>
A;Cross-references: UNIPROT:P21109; UNIPARC:UP10000156FB7; GB:M58021; GB:J05705; NID:9165; C;Superfamily: vertebrate rhodopsin
C;Keywords: G protein-coupled receptor; glycoprotein; membrane protein; neutrophil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      interleukin-8 receptor, high affinity - rabbit N;Alternate names: fMLP receptor C;Species: Oryctolagus cuniculus (domestic rabbit) C;Date: 22-Jan-1993 #sequence_revision 22-Jan-1993 C;Accession: A23669
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A;Experimental source: neutrophils
A;Note: sequence extracted from NCBI backbone (NCBIN:81526, NCBIP:81530)
C;Superiamily: vertebrate rhodopsin
C;Keywords: G protein-coupled receptor; transmembrane protein
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                                                                                                                                                      LSILWPIWYHCRRPRYLSSVMCVLLWALSILEKMFCDFLFSGANS--VWCETSDFI 179
                                                                                                                                                                                                                                                                                            GFLHSCLNPITYAFIG----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 YLLNLAMADLLF----ALTMPIWAVSKEKGWIFGTPLCKVVSLVKEVNFYSGILLLACIS
    FLLCGLPFG----
                                                                            TIAWLVFLCVVL-----CGSSLVLLVRILC-GSRKMPLTRLYV-----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TGMPPVE-KDYSPCLVVTQTLNKYVVVVIYALVFLLSLLGNSLVMLVILYSRSNRSVTDV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TKLTPINGREETPC--YNOTLS---FTGLTCIISLVALTGNA-VVLWLLGCRMRRNAVSI
                                        TAKW----CMVLRILPHTFGFILPLLVMLFCYGFTLRTLFQAHMGQKHRAMRVIFAVVLI
                                                                                                                  LAIVOST-RTLTQKRHLVKFICLGIWALSLILSLPFFLFRQ-VFSPNNSSPVCYEDLGHN 198
                                                                                                                                                                                            YLLNLAMAP-AFCPDHAYLGRLQGKRLDFRTPLCKVVSLVKEVNFYSGILLLACISVDRY 140
                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                   14.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14.0%; Score 237; DB 2; 27.6%; Pred. No. 3.4e-12.
                                                                                                                                                                                                                                                                                                                                                 54;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      60; Mismatches 104;
                                                                                                                                                                                                                                                                                                                                                                   Score 236.5; DB 2
Pred. No. 3.7e-12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ONFRNGFLKMLAARGL 330
    IQWALFSRIHLDWKVLFCHVHLVSIFLSALNSSANPII
                                                                                                                                                                                                                                                                                                                                                   Mismatches 117;
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                                                                              --TILLTVLV
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A;Cross-references: GDB:128855; OMIM:136539
A;Map position: 19q13.3-19q13.4
A;Introns: #status absent
C;Superfamily: vertebrate rhodopsin
C;Keywords: chemotaxis; G protein-coupled re
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: UNIPROT:P25089; UNIPARC:UPI0000050485; GB:M76673; NID:g182668; PII
C;Comment: This fMet-Leu-Phe receptor homolog, whose ligand is not yet known, appears
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R.Bao, L.; Gerard, N.P.; Eddy Jr., R.L.; Shows, T.B.; Genomics 13, 437-440, 1992
A;Title: Mapping of genes for the human C5a receptor (C: A;Reference number: A42009; MUID:92307681; PMID:1612600
A;Accession: C42009
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: DNA
A; Residues: 1-353 <BAO>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Status: nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change C;Accession: C42009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N;Alternate
C;Species: F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FMLP-related receptor 2 - human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Species: Homo sapiens (man)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         names: FMLP-related receptor I; probable chemotactic receptor FPRH2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                             LDRCICVLHPAWAQNHRTMSLAKRVMTGLWIFTIVLTLPNFIFWTTISTTNGDTYCIFNF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PINGREE-----TPCYNQTLSFTGLTCIISLVALTGNAVVLWLLGCRMRRNAVSIYI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     YAFIG----ONFRNGFLKMLAARGL 329
                                                                                                                                                                                                                                                                                                                                                 SRKMPLTRLYVTILLTVLVFLLCGLPF---GIQWALFSR---IHLDWKVLFCHVHLVSIF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TERCLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMF---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LNLALADFSFSA----ILPFRMVSVAMREKWPFASFLCKLVHVMIDINLFVSVYLITIIA 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LNLVAANFLFLSGHIIFSPLPLINI----RHP----ISKILSPVMTFPYFIGLSMLSAIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      YFFVGSFRQRQNRQN--LKLVLQRAL
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                                                                                                                                                                                                                                           LAFFNSCLNPILYVFMGRNFQERLIRSLPTSLERALTEVPD
                                                                                                                                                                                                                                                                               LSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPE 300
                                                                                                                                                                                                                                                                                                                  SR--PL-RVFAAV---VASFFICWFPYELIGILMAVWLKEMLLNGKYKIILVLINPTS-S
                                                                                                                                                                                                                                                                                                                                                                                       ----AFWGDTAVERLNVFITMAKVFLILHFIIGFTVPMSIITVCYGIIAAKIHRNHMIKS
                                                                                                                                                                                                                                                                                                                                                                                                                          SGANSVWCET----SDFITIAWLVFLCVVLCGSSLVLLVRILC--------G
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N.P.; Eddy Jr., R.L.; Shows, T.B.; Gerard,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       77;
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Pred. No. 1.9e-11;
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A;Status: nucleic acid sequence not shown
A;Molecule type: DNA
A;Residues: 1-369 <YAM>
A;Cross-references: UNIPROT:P30875; UNIPARC:UPI0000000A47; GB:M81832; NID:g201060; FA;Cross-references: UNIPROT:P30875; UNIPARC:UPI0000000A47; GB:M81832; NID:g201060; FR;Elliott, D.E.; Metwali, A.; Blum, A.M.; Sandor, M.; Lynch, R.; Weinstock, J.V.
J. Immunol. 153, 1180-1186, 1994
A;Title: T lymphocytes isolated from the hepatic granulomas of schistosome-infected A;Reference number: 156236; MUID:94300079; PMID:7913111
A;Accession: 156236
                                                                                                                                                                                                                                                                                                                                                 A;Title: Expression cloning of a rat brain so A;Reference number: A45291; MUID:92262491; PM A;Accession: A45291
A;Status: preliminary A;Molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Molecule type: mRNA
A;Residues: 99-309 <RES>
A;Cross-references: UNIPARC:UPI00001778F0; GB:S71756; NID:g560631
C;Superfamily: vertebrate rhodopsin
C;Keywords: G protein-coupled receptor; hormone receptor; transme
                                                                                                                                                                                                                                                                                                                                                                                                                                                     somatostatin receptor, somatotropin release-inhibiting factor receptor, SRIF receptor C;Species: Rattus norvegicus (Norway rat) (C;Date: 25-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004 C;Accession: A4591 C;Accession: A4591 R;Kluxen, F.W.; Bruns, C.; Lubbert, H. Proc. Natl. Acad. Sci. U.S.A. 89, 4618-4622, 1992
                                                                                                                                                                                                                         A;Molecule type: mRNA
A;Residues: 1-369 <KLU'>
A;Residues: 1-369 <KLU'>
A;Residues: 1-369 <KLU'>
A;Rocs-references: UNIPROT:P30680; UNIPARC:UPI0000135FF7; GB:M93273; NID:g207026; PIDN:I
A;Note: sequence extracted from NCBI backbone (NCBIN:102315, NCBIP:102316)
C;Superfamily: vertebrate rhodopsin
C;Keywords: G protein-coupled receptor; transmembrane protein
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Matches 94
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Best Local S
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Local Similarity 28.0%; Pred. No. 5.5e-11;
1es 94; Conservative 58; Mismatches 129;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               253
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                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RCLSILWPI----WYHCRRPRYLSSVMCVLLWALSLLRSILEWMF-------
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LNGSLGPSNGSNQTEPYYDMTSNAVLTFIYFVVCVVGLCGNTLVIYVILRYAKMKTITNI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               KVTRMVSIVVAVFIFCWLPFYIFNVSSVSVAISPTPAL--KGMFDFV----VILTYANSC 306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     YVT--ILLTVLVFLLCGLPF-----GIQWALFSRIHLDWKVLFCHVHLVSIFLSALNSS 266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INWPGESGAW--YTGFIIYAFILGFLVPLTIICLCYLFIIIKVKSSGIRVGSSKRKKSEK 252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LGTKLTPINGREET-PCYNOTLS--FTGLTCIISLVALTGNAVVLW-LLGCRMRRNAVSI
                                                                                    LGTKLTPINGREET-PCYNQTLS--FTGLTCIISLVALTGNAVVLW-LLGCRMRRNAVSI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   YILNLAIADELFMLG-LPFLAMQVALVHWPFGKAICRVVMTVDGINQFTSIFCLTVMSID
  YILNLVAANFLFLSGHIIFSPLPLINIRHPISK-ILSPVMT---FPYFIGLSMLSAISTE
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                                            LNGSLGPSNGSNQTEPYYDMTSNAVLTFIYFVVCVVGLCGNTLVIYVILRYAKMKTITNI
                                                                                                                                       Conservative
                                                                                                                                       13.2%; Score 222.5; DB 2; 28.0%; Pred. No. 5.5e-11; tive 58; Mismatches 129;
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PMID:1374909
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E.; Navarro

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A;Molecule type: mRNA
A;Residues: 1-346 </AN>
A;Residues: 1-346 </ANS
A;Cross-references: UNIPROT:P30875; UNIPARC:UPI0000044D82; EME
C;Superfamily: vertebrate rhodopsin
C;Keywords: G protein-coupled receptor; transmembrane protein
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S29248
interleukin-8 receptor (clone 5Bla) - rabbit
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995
C;Accession: A53752
                                                                       RESULT
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A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Title: Cloning and expression of a novel mouse somatostatin receptor (SSTR2B) A; Reference number: S29248; MUID:93012001; PMID:1397330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R; Vanetti, M.; Kouba, M.; Wang, X.; Vogt, G.; Hoellt, FEBS Lett. 311, 290-294, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    somatostatin receptor 2B - mouse
C;Species: Mus musculus (house mouse)
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 09-Jul-2004
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                                                                                                                                                                                                                                                 YVT--ILLTVLVFLLCGLPF-----GIQWALFSRIHLDWKVLFCHVHLVSIFLSALNSS
                                                                                                                                                                                                                                                                                                                   FLFSGANSVWCETSDFITIAWLVFLCV---VLCGSSLVLLVR-----ILCGSRKMPLTRL
                                                                                                                                                                                                                                                                                                                                                                                                                                                             YILNLVAANFLELSGHIIFSPLPLINIRHPISK-ILSPVMT---FPYFIGLSMLSAISTE
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                                                                                                                                                                                                              KVTRMVSIVVAVFIFCWLPFYIFNVSSVSVAISPTPAL-
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                                                                                                                                            -DNFKKSFQNVL 327
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Pred. No. 1.:
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                                                                                                                                                                                                              -KGMFDFV----VILTYANTC
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R;Prado, G.N.; Thomas, K.M.; Suzuki, H.; LaRosa, G.J.; Wilkinson, N.; Folco, E.; Na
J. Biol. Chem. 269, 12391-12394, 1994
A;Title: Molecular characterization of a novel rabbit interleukin-8 receptor isotyp
A;Reference number: A53752; MUID:94230294; PMID:8175642
A;Accession: A53752
A;Status: preliminary
A;Molecula type: mRNA
A;Residues: 1-358 <PRA>
A;Residues: 1-358 <PRA>
A;Residues: 1-358 <PRA>
A;Residues: 1-358 <PRA>
A;Cross-references: UNIPROT:935344; UNIPARC:UPI000012D4F4; GB:L24445; NID:g437661;
C;Superfamily: vertebrate rhodopsin
C;Keywords: G protein-coupled receptor; transmembrane protein
                                                                                                                                                                     A;Status: preliminary
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-15 <RE2>
A;Cross-references: UNIPARC:UPI00000053D; EMBL:U11872; NID:g511808; PIDN:AAA64380.1;
A;Cross-references: UNIPARC:UPI000000053D; EMBL:U11877; NID:g511818; PID:
14; PID:g511815; EMBL:U11876; NID:g511816; PID:g511817; EMBL:U11877; NID:g511818; PID:
R;Sprenger, H.; Lloyd, A.R.; Lautens, L.L.; Bonner, T.I.; Kelvin, D.J.
J. Biol. Chem. 269, 11065-11072, 1994
A;Title: Structure, genomic organization, and expression of the human interleukin-8 re
A;Reference number: A53611; MUID:94209273; PMID:7512557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             interleukin-8 receptor type B - human
(Species: Homo sapiens (man)
(Species: Homo sapiens (man)
(Date: 07-Oct-1994 #sequence_revision 12-Apr-1996 #text_change 09-Jul-2004
(CAccession: I37998; I38712; A53611; A39446
R;Ahuja, S.K.; Shetty, A.; Tiffany, H.L.; Murphy, P.M.
J. Biol. Chem. 269, 26381-26389, 1994
A;Title: Comparison of the genomic organization and promoter function for ha;Reference number: I37898; MUID:95014476; PMID:7929358
A;Accession: I37898
A;Cross-references: UNIPARC:UPI00000746D6; R;Murphy, P.M.; Tiffany, H.L. Science 253, 1280-1283, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-360 <RES>
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A; Status: preliminary
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                                                                         A; Residues: 6-360 <SPR>
                                                                                                      A; Molecule type: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TAKWRMVLRILPQTFGFILPLLVMLFCYVFTLRTLFQAHMGQKH---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       279
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -VLLTDTLMRTHVIQETCERRNDIDRALDATEILGFLHSCLNPI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       53;
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Pred. No. 1.7e-10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches 105;
                                                 GB:M99412; GB:L19593
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ----VSIFLSALNSSANPI
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                                                                                                                                                                                                                                                                                                                                                                                                                                    PID
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:95)
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N;Alternate names: somatotropin release-inhibiting factor (Species: Rattus norvegicus (Norway rat) (C;Species: O2-Aug-1996 #sequence revision O2-Aug-1996 #text_cl C;Date: O2-Aug-1996 #sequence revision O2-Aug-1996 #text_cl C;Accession: I57940; I57949; S39244 R;O'Carroll, A.M.; Lolait, S.J.; Konig, M.; Mahan, L.C. Mol. Pharmacol. 42, 939-946, 1992 Mol. Pharmacol. 42, 939-946, 1992 A;Title: Molecular cloning and expression of a pituitary so A;Reference number: I57940; MUID:93125499; PMID:1362243 A;Accession: I57940
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A,Title: Cloning of complementary DNA encoding a functional human A,Reference number: A39446; MUID:91368200; PMID:1891716 A,Accession: A39446
                                                                                        A;Status: preliminary; translated A;Molecule type: mRNA A;Residues: 341-363 <OCA2>
                                                                                                                                                         A;Cross-references: UNIPROT: P30938; UNIPARC: UPI0000135FFB; R;O'Carroll, A.M.; Lolait, S.J.; Konig, M.; Mahan, L.C. Mol. Pharmacol. 44, 1278, 1993
A;Title: Molecular cloning and expression of a pituitary su A;Reference number: 157949; MUID:94088493; PMID:8264565
A; Description:
                    A; Experimental source: pituitary R; Penetta, R.; Greenwood, M.; Patel, Y.C. submitted to the EMBL Data Library, Augus
                                                                                                                                            A; Accession: I57949
                                                                                                                                                                                                                                                  A; Molecule type: mRNA
A; Residues: 1-363 < OCA1>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C;Genetics:
A;Gene: GDB:IL8RB; IL8RA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Cross-references: UNIPARC:UPI00000746D6; GB:M73969 C;Comment: This receptor, unlike ILBRA, binds severa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Status: preliminary; nucleic A;Molecule type: mRNA A;Residues: 6-360 <MUR>
                                                                                                                                                                                                                                                                                      A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                somatostatin receptor 5 - rat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;Superfamily: vertebrate rhodopsin;Keywords: G protein-coupled recept
                                                                        Cross-references: UNIPARC:UPI00000004EB; GB:S67370; NID:g455947; PIDN:AAB29371.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   95 LLFALTLPIWAASKVNGWIFGTFL------CKVVSLLKEVNFYSGILLLACISVDRY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              81;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FFVG-SFR 279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LSILWPIWYHCRR----PRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWC-ETS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DAAPCEPESLEINKYFVVIIYALVFLLSLLGNSLVMLVILYSRVGRSVTDVYLLNLALAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EETPCYNQTLSFTG-----LTCIISLVALTGNA-VVLWLLGCRMRRNAVSIYILNLVAAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein-coupled receptor; transmembrane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AFIGQKFR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GNNTANWRMLLRILPQSFGFIVPLLIMLFCYGFTLRTLFKAHMGQKHRAMRVIFAVVLIF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DFITIAWLVFLCVV--LCGSSLVLLVRILC-GSRKMPLTRLYV-----TILLTVLVF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LAIV----HATRTLTQKRYLVKFICLSIWGLSLLLALPVLLFRRTVYSSNVSPACYEDM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LLCWLPYNLVLLADTLMRTQVIQETCERRNHIDRALDATEI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FLF------LSGHIFSPLPLINIRHPISKILSPVMTFPYFIGLSMLSAISTERC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          55;
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     nucleotide
                                                                                                                              from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    - IQWALFSRIHLDWKVLFCHVHLVSIFLSALNSSANPIIY
                    August 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                            GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             215.5; DB
No. 2e-10;
     and
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     amino
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 acid sequence
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                                                                                                                                                                                   somatostatin
                                                                                                                                                                                                                                                                                                                                               somatostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                     subtype 28 receptor
                                                                                                                                                                                                                                     GB:L04535; NID:G409238;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LGILHSCLNPLIY
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                                                                                                                                                                                                                                                                                                                                                                                                                    09-Jul-2004
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   0f
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              interleukin-8,
     the
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     rat
     somatosta
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A;Reference number: S39244
A;Accession: S39244
A;Molecule type: mRNA
A;Residues: 309-363 <PEN>
A;Cross-references: UNIPARC:UPI00001707D8;
C;Genetics:
A;Gene: SSTR5
A;Gene: SSTR5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
314
                                       296
                                                                                                                                                             210
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  ELSG 317
                                                                                                                                                                                                                                        L--FSGANSVWCETSDFITIAWLVFLCV-----VLCGSSLVLLVRILCGSRKM----
                                                                                                                                                                                                                                                                                TSIFCLWVMSVDRYLAVVHPLRSARW---RRPR-VAKWASAAVWVFSLLMSLPLLVFADV 178
                                                                                                                                                                                                                                                                                                                      IGLSMLSAISTERCLSILWPI----WYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDF 163
                                                                                                                                                                                                                                                                                                                                                                                                      RNAVSIYILNLVAANFLELSGHIIFSPLPLINIRHPI----SKILSPVMTFP---YF
                                                                             NSSANPIIYFFVG-SFRORONRONLKLVLOR--ALODT----PEVDEGG----GWLPOETL
                                                                                                                       RRRSEPKVTRMVVVV----VLVFVGCWLPFFIVNIVNLAFTLPEEPTSAGLYFFVVVLSYA
                                                                                                                                                                                                     QEGWGTCNLSWPEPVGLWGAAFITYTSVLGFFGPLLVICLCYLLIVVKVKAAGMRVGSSR
                                                                                                                                                                                                                                                                                                                                                                  KTVTNVYILNLAVADVLFMLG-----LPFLATQNAVVSYWPFGSFLCRLVMTLDGINQF
                                         NSCANPLLYGELSDNFROSFRK----VLCLRRGYGMEDADAIEPRPDKSGRPOATLPTRSC 352
                                                                                                                                                                                                                                                                                                                                                                                                                                                TPSWNASAASSGNHNWSLVGSASPMGARAVLVPVLYLLVCTVGLSGNTLVIYVVLRHAKM
                                                                                                                                                             ----PLTRLYVTILLTVLVFLLCGLPFGIQWALFSRIHLDWKVLFCHVHLVSIFLSAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12.7%; Score 215.5; DB 2
24.5%; Pred. No. 2.1e-10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  66;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL: X74828; NID: g433911; PIDN: CAA52825.1; PIL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ---LTCIISLVALTGNAVVLW-LLGCRMR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  126;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  83;
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                                                                                                                                                               263
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                                                                                                                                                                                                                                               209
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Gene 118, 303-304, 1992
A;Title: Cloning of a cDNA encoding a receptor related to the formyl peptide a A;Title: Cloning of a cDNA encoding a receptor related to the formyl peptide a A;Reference number: JC1258; MUID:92380523; PMID:1511907
A;Accession: JC1258
A;Atatus: nucleic acid sequence not shown
A;Atatus: nucleic acid sequence not shown
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-351 < PER>
A;Cross-references: UNITARC: UPI0000012D0; EMBL:X63819; NID:g31460; PID:g31461
A;Experimental source: bone marrow mRNA
A;Experimental source: bone marrow m Genomics 13, 437-440, 1992

A;Title: Mapping of genes for the human C5a receptor (C5AR), human FV A;Reference number: A42009; MUID:92307681; PMID:1612600

A;Accession: B42009

A;Accession: B42009

A;Status: nucleic acid sequence not shown

A;Holecule type: DNA

A;Residues: 1-263,'A',265-338,'C',340-351 <BAO>

A;Residues: 1-263,'A',265-338,'C',340-351 <BAO>

A;Cross-references: UNIPROT:P25909; UNIPRACTIVIDIO001778D8; GB:M76672

A;Hote: authors translated the codons GTG for residue 15 as G1u, TCT

R;Perez, H.D.; Holmes, R.; Kelly, E.; McClary, J.; Andrews, W.H.

Gene 118, 303-304, 1992 N;Alternate names: FMLP receptor homolog FPR2; formyl peptide receptor like-1; probable C;Species: Homo sapiens (man) C;Date: 30-Sep-193 #sequence revision 14-Jul-1995 #text change 09-Jul-2004 C;Accession: B42009; JC1258; JQ1521; A42492; I54751; S21581 C;Accession: B42009; JC1258; JQ1521; A42492; I54751; S21581 C;Accession: B42009; JC1258; JQ1521; A42492; I54751; S21581 FMLP-related receptor 1 - human FMLP PID: 931461 for receptor (FPR), 19 œ.

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peptide

receptor

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A;Cross-references: GDB:127554; OMIM:136538
A;Map position: 19q13.3-19q13.4
A;Introns: #status absent
C;Superfamily: vertebrate rhodopsin
C;Keywords: chemotaxis; G protein-coupled receptor; glycoprotein; transmembrane #status predicted <TM1>
F;27-53/Domain: transmembrane #status predicted <TM2>
F;100-121/Domain: transmembrane #status predicted <TM3>
F;100-121/Domain: transmembrane #status predicted <TM3>
F;106-226/Domain: transmembrane #status predicted <TM5>
F;206-226/Domain: transmembrane #status predicted <TM6>
F;242-266/Domain: transmembrane #status predicted <TM6>
F;242-266/Domain: transmembrane #status predicted <TM7>
F;44/Binding site: carbohydrate (Asn) (covalent) #status predicted
F;98-176/Disulfide bonds: #status predicted
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A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-351 <RES>
A;Cross-references: UNIPARC:UPI0000012D0; GB:D10922; NID:g219864; PIDN:BAA01720.1;
A;Cross-references: UNIPARC:UPI00000012D0; homologous to the N-formyl peptide recept
C;Comment: This G-protein coupled receptor, homologous to the N-formyl peptide recept
differentiated myeloid cells and is probably a chemotactic receptor for some other l
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R;Nomura, H.; Nielsen, B.W.; Matsushima, K.
Int. Immunol. 5, 1239-1249, 1993
Int. Immunol. 5, 1239-1249, 1993
A;Title: Molecular cloning of cDNAs encoding a LD78 receptor and putative leukocyte chem A;Reference number: 154751; MUID:94092629; PMID:7505609
A;Accession: I54751
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A;Note: formyl peptide stimulated calcium mobilization comparable to that of the formyl
R;Nurphy, P.M.; Ozcelik, T.; Kenney, R.T.; Tiffany, H.L.; McDermott, D.; Francke, U.
J. Biol. Chem. 267, 7637-7643, 1992
A;Title: A structural homologue of the N-formyl peptide receptor. Characterization and calcium an
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A; Residues: 1-351 < MUR>
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     NPTSSLAFFNSCLNPMLYVFVGQDFRERL-IHSLPTSLERALSE
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Search completed: February

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Job time : 44 secs

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## **Protein Sequence Searches - February 2005**

All of the sequence databases on ABSS have recently been updated.

- Please note that the curators of the UniProt database have purged some temporary accession numbers from the most recent version of UniProt. These sequences have been assigned new permanent accession numbers. The new UniProt record may not contain the previous temporary accession number.
- If you encounter an accession number from an older search run against UniProt (results file extension .rup) that can no longer be found in the database, the permanent record with the new accession number can be found by searching the old accession number in the UniProt Protein Archive database (UniPARC) at:

http://www.pir.uniprot.org/database/archive.shtml

If you have any questions regarding this information or your results, please contact any STIC searcher.

When submitting sequence search results for scanning into IFW, please include a copy of this attachment to assist any future Examiners or members of the public who may encounter UniProt temporary accession numbers.

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Minimum DB
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L.H., Shen S.I., Rub Cas C., Abr Male Chenko Chenko Myee	ubMed=11551509; DOI=10.1016/ Zylka M.J., Simon M.I., And F GPCRs expressed in specifi neurons."; O1).  VARIANT ASN-169, AND TISSU ubMed=11850634; DOI=10.1038/ ini E., Groblewski T., O'Dor ini E., Schmidt R., Pelle Y., Banville D., Shen S., & Ahmad S.; ne products activate a new f Rs."; 1-209(2002).	IT 1 3 HUMAN 7 HUMAN STANDARD; PRT; 3 Q96LB0; Q8TDE1; 25-OCT-2004 (Rel. 45, Created) 25-OCT-2004 (Rel. 45, Last sequence up 13-SEP-2005 (Rel. 48, Last annotation Mas-related G-protein coupled receptor specific G-protein coupled receptor 1) Name=MRGPRX3; Synonyms=MRGX3, SNSR1; Homo sapiens (Human). Bukaryota; Metazoa; Chordata; Craniata Mammalia; Eutheria; Euarchontoglires; Homo.	32 773 45.7 331 2 Q5FVUI RAT 33 765 45.2 304 1 MRGA1 MOUSE 34 749 44.3 338 2 Q8CDY4 MOUSE 35 749 44.3 338 2 Q91ZC2 MOUSE 36 744 44.0 294 2 Q7TN48 RAT 37 730 43.2 302 1 MRGA3 MOUSE 38 729.5 43.1 301 1 MRGA6 MOUSE 42.7 305 1 MRGA7 MOUSE 42.7 305 1 MRGA7 MOUSE 41.6 338 2 Q91ZC3 MOUSE 41 704 41.6 338 2 Q91ZC3 MOUSE 41 704 41.6 338 2 Q91ZC3 MOUSE 42 685.5 40.5 313 1 MRGA5 MOUSE 43 680 40.2 304 1 MRGA5 MOUSE 44 677.5 40.1 323 2 Q7TN45 RAT 45 675.5 39.9 305 1 MRGA2_MOUSE 45 675.5 39.9 305 1 MRGA2_MOUSE
O73/pnas.242603899; Derge J.G., Berge J.G., Inmen C.M., Schuler G.D., aefer C.F., Bhat N.K., Wang J., Hsieh F., bin G.M., Hong L., savant T.L., Scheetz T.E., Carninci P., Prange C., ramson R.D., Mullahy S.J., ek J.A., Gunaratne P.H., ek J.A., Gunaratne P.H., k.J.A., Gy L.J., Hulyk S.W., Lu X., Gibbs R.A., Lu X., Gibbs R.A., Du X., Gibbs R.A., o Y., Bouffard G.G., o Y., Bouffard G.G., o Y., Smailus D.E., a U., Smailus D.E., a U., Smailus D.E.,	S0092-8674(01)00483-4; lerson D.J.; c subsets of f SPECIFICITY. fnn815; nne11 D., Roy MO., trier M., Labarre M., stroem P., Payza K., stroem P., Payza K.,	22 AA. ddate) update) . member X3 (Sensory neuron- . Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae;	Q5fvul rattus norv Q91ww5 mus musculu Q8cdy4 mus musculu Q91zc2 mus musculu Q7tn48 rattus norv Q91ww3 mus musculu Q91zc6 mus musculu Q91zc6 mus musculu Q91zc3 mus musculu Q91zc3 mus musculu Q91zc3 mus musculu Q91zc7 mus musculu Q91xc7 mus musculu

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"Generation and initial analysis of more than 15,000 full-length human

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PRINTS; PR00237; GPCREIN RECEP F1 1; 1.

PROSITE; PS00237; GPROTEIN RECEP F1 2; 1.

PROSITE; PS50262; GPROTEIN RECEP F1 2; 1.

G-protein coupled receptor; Polymorphism; Receptor; Transducer;
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EMBL; AF474987; AAL86878.2; -; GENOMIC_DNA.
EMBL; BC067292; AAH67292-1; -; MENUA.
Ensembl; ENSG00000179826; Homo sapiens.
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C. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).

FUNCTION: Orphan receptor. Probably involved in the function of nociceptive neurons. May regulate nociceptor function and/or development, including the sensation or modulation of pain.

Potently activated by enkephalins (By similarity).

SUBCELLULAR LOCATION: Integral membrane protein.

TISSUE SPECIFICITY: Uniquely localized in a subset of small dorsal root and trigeninal sensory neurons.

SIMILARITY: Belongs to the G-protein coupled receptor 1 family.
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MEDILINE=21853733; PubMed=11850634; DOI=10.1038/nn815;
Lembo P.M.C., Grazzini E., Groblewski T., O'Donnell D., I
Zhang J., Hoffert C., Cao J., Schmidt R., Pelletier M., 1
Gosselin M., Fortin Y., Banville D., Shen S., Stroem P.,
Dray A., Walker P., Ahmad S.;
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prosite; ps50262; G_PROTEIN_RECEP_F1_2; 1.
G-protein coupled receptor; Glycoprotein; Receptor; Transducer;
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Ensembl; ENSG0000179826; Homo Bapiens.
InterPro; IPR000276; GPCR_Rhodpsn.
Pfam; PF00001; 7tm_1; 1.
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Neurosci. 5:201-209(2002).

FUNCTION: Orphan receptor. Probably involved in the function rociceptive neurons. May regulate nociceptor function and/or development, including the sensation or modulation of pain. Potently activated by enkephalins (By similarity).

SUBCELLULAR LOCATION: Integral membrane protein.

TISSUE SPECIFICITY: Uniquely localized in a subset of small
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                                                                                  They are the first and the first activate a new family of sensory to neuron-specific GPCRe."

What. Neurosci. 5:201-209(2002).

Nat. Neurosci. 5:201-209(2002).

C-!-FUNCTION: Orphan receptor. Probably involved in the function of nociceptive neurons. May regulate nociceptor function and/or development, including the sensation or modulation of pain.

CC notently activated by enkephalins including BAM22 (bovine adrenal medulla peptide 22) and BAM (8-22). BAM22 is the most potent compound and evoked a large and dose-dependent release of intracellular calcium in stably transfected cells. G(alpha)q

CC intracellular calcium in stably transfected cells. G(alpha)q

CC root and trigeminal sensory neurons.

CC -I-SIMILARITY: Belongs to the G-protein coupled receptor 1 family.
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Best Local S
Matches 302
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25-OCT-2004 (Rel. 45, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Sensory neuron-specific G-protein and Name-SNSR3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3 HUMAN
SNSR3 HI
QBTDD9;
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CARBOHYD
SEQUENCE
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MEDLING=21853733; PubMed=11850634; DOI=10.1038/nn815;
Lembo P.M.C., Grazzini E., Grobhewski T., O'Donnell D., Roy M.-C
Zhang J., Hoffert C., Cao J., Schmidt R., Pelletier M., Labarre
Gosselin M., Fortin Y., Banville D., Shen S., Stroem P., Payza K
Dray A., Walker P., Ahmad S.;
                            This
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=9606;
  Swiss-Prot en
een the Swiss
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02; Conservative
                                                                             subfamily.
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     Institute
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  copyright. It is productive of Bioinformatics
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N-linked (GlcNAc. . .) (P
DBC24308E3B4611B CRC64;
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  produced through a catics and the EMBL
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                       a collaboration
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Matches 267
MRGX1_HUMAN STANDARD; PRT; 322 AA.
Q96LB2; QBTDDB;
Q96LB2; QBTDDB;
25-OCT-2004 (Rel. 45, Created)
25-OCT-2004 (Rel. 45, Last sequence update)
13-SEP-2005 (Rel. 48, Last annotation update)
13-SEP-2005 (Rel. 48, Last annotation update)
Mas-related G-protein coupled receptor member X1
specific G-protein coupled receptor 4).
Name=MRGPRX1, Synonyms=MRGX1, SNSR4;
Homo sapiens (Human).
Bukaryota; Metazoa; Chordata; Craniata; Vertebra
Eukaryota; Metazoa; Surdata; Craniata; Primates;
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PROSITE; PS00237; G_PROTEIN RECEP_F1_1; 1.

PROSITE; PS50262; G_PROTEIN RECEP_F1_2; 1.

G-protein coupled receptor; Glycoprotein; Receptor; Transducer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; AF474989; AAL86880.2; -; Ger
Ensembl; ENSG00000170255; Homo sar
InterPro; IPR000276; GPCR_Rhodpsn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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82.9%; Pred
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4 (Potential).

Extracellular (Potential).

5 (Potential).

Cytoplasmic (Potential).

6 (Potential).

Extracellular (Potential).

Extracellular (Potential).

Cytoplasmic (Potential).

7 (Potential).

Cytoplasmic (Potential).

Cytoplasmic (Potential).

N-linked (GlcMac. .) (Potential).
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1 (Potential).

Cytoplasmic (Potential).

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No. 1.9e-92;
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Chordata; Craniata; Vertebrata; Euarchontoglires; Primates; Cata

Catarrhini; Hominidae;

Euteleostomi;

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(Sensory

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G-protein co
Transducer;
TOPO_DOM
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MEDLINE=21853733; PubMed=11850634; DOI=10.1038/nn815;
Lembo P.M.C., Grazzini E., Groblewski T., O'Donnell D., I
Zhang J., Hoffert C., Cao J., Schmidt R., Pelletier M., I
Gosselin M., Fortin Y., Banville D., Shen S., Stroem P.,
Dray A., Walker P., Ahmad S.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=22040266; PubMed=12044878; DOI=10.1016/S0014-5793(02)02775-8; Takeda S., Kadowaki S., Haga T., Takaesu H., Mitaku S.; "Identification of G protein-coupled receptor genes from the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nociceptive sensory neurons."; Cell 106:619-632(2001).
              TRANSMEM
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                                                                                                       Pfam; PF00001; 7tm 1; 1.

PRINTS; PR00237; GFCRRHODOPSN.

PROSITE; PS00237; G PROTEIN RE

PROSITE; PS50262; G PROTEIN RE

G-protein coupled receptor; Gl
                                                                                                                                                                                                                                                                                                                                                                                   This
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NCBI_TaxID=9606;
                                                                                                                                                                                                           Ensembl; ENSG00000170255;
HGNC; HGNC:17962; MRGPRX1.
                                                                                                                                                                                                                                               EMBL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Suwa M., Sato T., Okouchi I., Arita M., Futami K., Tsutsumi S., Aburatani H., Asai K., Akiyama Y.; "Genome-wide discovery and analysis of human seven
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Suwa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=21435808;
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                                                                                                                                                                                 nterPro; IPR000276; GPCR_Rhodpsn.
                                                                                                                                                                                                                                                                                                                                                                                                                         FUNCTION: Orphan receptor. Probably involved in the function of modiceptive neurons. May regulate nociceptor function and/or development, including the sensation or modulation of pain. Potently activated by enkephalins including BAM22 (bovine adrenal medulla peptide 22) and BAM (8-22). BAM22 is the most potent compound and evoked a large and dose-dependent release of intracellular calcium in stably transfected cells. G(alpha) q proteins are involved in the calcium-signaling pathway. SUBCELIULAR LOCATION: Integral membrane protein.

TISSUE SPECIFICITY: Uniquely localized in a subset of small dorsal root and trigeninal sensory neurons.
SIMILARITY: Belongs to the G-protein coupled receptor 1 family.
                                                                                                                                                                                                                                                                                                                                                     European
                                                                                                                                                                                                                                                                                                                                                                                                                 Mas subfamily.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FUNCTION: Orphan receptor.
                                                                                                                                                                                                                                                                                                                               Swiss-Prot entry is copyright. It is proude the Swiss Institute of Bioinformatics een the Swiss Institute of There a European Bioinformatics Institute. There a
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AF474990; AAL86881.1; -;
AB083628; BAB89341.1; -;
AB065846; BAC06064.1; -;
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435808; PubMed=11551509; DOI=10.1016/S0092-8674(01)00483-4;
435808; PubMed=11551509; DOI=10.1016/S0092-8674(01)00483-4;
435808; PubMed=11551509; DOI=10.1016/S0092-8674(01)00483-4;
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 Transmembrane.
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_RECEP_F1_2; 1.
_Glycoprotein; Polymorphism;
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Cytoplasmic (Po
2 (Potential).
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Matches 264
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25-OCT-2004
25-OCT-2004
10-MAY-2005
                                                                                                                               NUCLEOTIDE SEQUENCE, AND TISSUE SPECIFICITY.

MEDLINE=21853733; PubMed=11850634; DOI=10.1038/nn815;

Lembo P.M.C., Grazzini E., Groblewski T., O'Donnell D., J

Zhang J., Hoffert C., Cao J., Schmidt R., Pelletier M., J

Gosselin M., Fortin Y., Banville D., Shen S., Stroem P.,

Dray A., Walker P., Ahmad S.;
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25-OCT-2004 (Rel. 45, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation updat
Sensory neuron-specific G-protein coupled x
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rimates; Catarrhini; Hominidae;
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Labarre M.,
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Hominidae;

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RESULT 6
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Q4V912 HUMAN PRELIMINARY;
ID Q4V912;
AC Q4V912;
DT 13-SEP-2005 (TrEMBLrel. 3
DT 13-SEP-2005 (TrEMBLrel. 3
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PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
G-protein coupled receptor; Glycoprotein; Receptor; Transducer;
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Ensembl; ENSG00000179817; Homo sapiens.
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LARITY: Belongs to the G-protein coupled receptor 1 family.
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5 (Potential).
5 (Potential).
6 (Potential).
6 (Potential).
Extracellular (Potential).
7 (Potential)
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1 (Potential).
Cytoplasmic (Potential).
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Extracellular (Potential).
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N-linked (GlcNAc...) (Potentiai)
3D6FFB4B5DDFDD90 CRC64;
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Pred. No. 3
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     sequence update)
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RX MEDININE=22388557; pubMed=12477932; DOI=10.1073/pnas.242603899;

RX MEDININE=22388557; pubMed=12477932; DOI=10.1073/pnas.242603899;

RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

RA Klausner R.D., Collins F.S., Wagner L., Schammen C.M., Schuler G.D.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschak S.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gibbs R.A.,

RA Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

RA Millian M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

RA Molting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

RA Molting W.R., Touchman J.W., Green E.D., Dickson M.C.,

RA Rodriguez A.C., Grimwood J.W., Schmutz J., Myers R.M.,

RGnerst G. M., Schmit J.E., Jones S.J.M., Marra M.A.;

RGnerst G. M., Schmit J.E., Jones S.J.M., Marra M.A.;

RGnerst G. G., Schmit J.E., Jones S.J.M., Marra M.A.;
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PRINTS; PR00237; GPCRRHODDPSN.

PROSITE; PS00237; G PROTEIN RECEP F1 1; UNKNOWN 1.

PROSITE; PS00263; G PROTEIN RECEP F1 2; 1.

G-protein coupled receptor; Receptor; Transducer; Transmembrane.

SEQUENCE 322 AA; 36236 MW; C7F219F4418E8AD1 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Homin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (JUN-2005) to the EMBL/GenBank/DDBJ databases.
-!- SUBCELLULAR LOCATION: Integral membrane protein (By similarity).
EMBL; BC096835; AAH96835.1; -; mRNA.
InterPro; IPRO10276; GPCR_Rhodpsn.
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   VDEGGGWLPQETLELSGSKLEQ
                                                                            RIHLDWKVLFCHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPE
                                                                                                                                                                                                CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMPCDFLFSGANSVWCETSDFIT
                                                                                                                                                                                                                                                                      FSIYILNLAAADFLFLSGRLIYSLLSFISIPHTISKILYPVMMFSYFAGLSFLSAVSTER
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                                                          WIHVDREVLECHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDASE
                                                                                                                           VAWLIFICVVLCGSSLVLLIRILCGSRKIPLTRLYVTILLTVLVFLLCGLPFGIQFFLFL
                                                                                                                                             IAMLVFLCVVLCGSSLVLLVRILCGSRKMPLTRLYVTILLTVLVFLLCGLPFGIQWALFS
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81.7%; Pred. No. 3.2¢
tive 23; Mismatches
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VDEGGGQLPEEILELSGSRLDQ 322

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RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RX Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gubs R.A.,
RA Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
"Generation and initial analysis of more than 15,000 full-length human pard and mouse cDNA accumences.":
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PROXA HUMAN STANDARD; PRT; 322 AA.

Q96LA9; Q502W0; Q8TDD6;
25-OCT-2004 (Rel. 45, Created)
25-OCT-2004 (Rel. 45, Last sequence update)
13-SEP-2005 (Rel. 48, Last annotation update)
13-SEP-2005 (Rel. 48, Last annotation update)
Mas-related G-protein coupled receptor member X4 (Sensory neuron-
specific G-protein coupled receptor 6).

Name=MRGPRX4; Synonyms=WRGX4, SNSR6;

Homo spaiens (Human). Charasta. Cranista: Vertebrata: Euteleostomi;
                                                                                         the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and mouse cDNA sequences."; Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
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NUCLEOTIDE SEQUENCE [LARGE SCALE MRNA].
NUCLEOTIDE SEQUENCE [LARGE SCALE MRNA].
242603899;
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"A diverse family of GPCRs expressed in specific subsets of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                            This
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MEDLINE=21435808; PubMed=11551509;
                                                                                                                                                                                                                                                                 FUNCTION: Orphan receptor. Probably involved in the function of nociceptive neurons. May regulate nociceptor function and/or development, including the sensation or modulation of pain. Potently activated by enkephalins (By similarity). SUBCELLULAR LOCATION: Integral membrane protein. TISSUE SPECIFICITY: Uniquely localized in a subset of small dorsal root and trigeminal sensory neurons.

SIMILARITY: Belongs to the G-protein coupled receptor 1 family.
                                          s Swiss-Prot entry is copyright. It is produced through a collabo
ween the Swiss Institute of Bioinformatics and the EMBL outste-
Buropean Bioinformatics Institute. There are no restrictions of
as long as its content is in no way modified and this statement
                                                                                                                                                                                                                                      Mas subfamily.
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Zylka M.J., Simon M.I., Anderson D.J.;
                                                                                                                                a collaboration -
MBL outstation -
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                                                                                                                                                                                                                                                                      VSIYILNLVAANFLFLSGHIIFSPLFLINIRHPISKILSPVMTFFYFIGLSMLSAISTER
                                                                                                                                                                                                                                                                                                                 PS00237; G_PROTEIN_RECEP_F1_1; 1.
PS50262; G_PROTEIN_RECEP_F1_2; 1.
n coupled receptor; Glycoprotein; Polymorphism; Receptor;
                                                           IAWLVFLCVVLCGSSLVLLVRILCGSRKWPLTRLYVTILLTVLVFLLCGLPFGIQWALFS
                                                                                                                                                                                                                                                    VSIYILNLAAADFLFLSFQIIRSPLRLINISHLIRKILVSVWTFPYFTGLSMLSAISTER
                                                                                                                        VAWLIFLCVVLCVSSLVLLVRILCGSRKMPLTRLYVTILLTVLVFLLCGLPFGILGALIY
 VDKGEGQLPEESLELSGSRL
                            VDEGGGWLPQETLELSGSKL
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                                                                                                                                                                                                                                                                                                                                                                                                   Score 1366; DB 1;
Pred. No. 6.1e-90;
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S -> L (in dbSNP:2445179)
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R -> K (in Ref. ;
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Q5U9EO_MACFA
ID Q5U9EO, WACFA PRELIMINARY;
AC Q5U9EO;
AC Q5U9EO;
DT 01-FEB-2005 (TrEMBLrel. 29,
DE Mas-related protein X1/7.
GN MGCAEO factorial (Crabe of Eukaryota, Metazoa; Chordat
OC Eukaryota, Metazoa; Chordat
OC Eukaryota, Metazoa; Chordat
OC Eukaryota, Metazoa; Chordat
OC Cercopithecidae; Cercopithe
OX NCBI_TAXID=9541;
RN NUCLEOTIDE SEQUENCE.
RA Zhang L., Taylor N., Ford R
RT "Cloning and Expression of
RT Human.";
RA Zhang L., Taylor N., Ford
RT Human.";
RT SUBCELLULAR LOCATION: I
DR GO; GO:00016021; C:integral
DR GO; GO:00016021; C:integral
DR GO; GO:00016021; C:integral
DR GO; GO:00016021; C:integral
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DR GO; GO:0007165; P:sdgnal tr
DR GO; GO:0007165; P:sdgnal tr
DR FANNTS; PR00237; G-PROTEIN
DR PROSITE; PS00237; G-PROTEIN
DR SEQUENCE 322 AA; 36781 M
RESULT 9
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OS Macaca fascicu
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Best Local S
Matches 255
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Name=MrgX8;
Macaca fascicularis
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Butheria; Euarchontoglires; Primates; Catarrhini;

Cercopithecidae; Cercopithecinae; Macaca.
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Last sequence
Last annotation
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Pred. No. 1.2e-
27; Mismatches
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Last annotation update)
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in Macaque, Mouse,
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Matches 244
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Q4QXU4;
13-SEP-2005 (7
13-SEP-2005 (7
13-SEP-2005 (7
                                                                                                                                                                             Trachypithecus francoisi (Francois' langur) (Indochinese langur). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Cercopithecidae; Colobinae; Trachypithecus.
NCBI_TaxID=54180;
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
Cercopithecidae; Cercopithecinae; Macaca.
NCBI_TaxID=9541;
NUCLEOTIDE SEQUENCE.
PubMed=15862286; DOI=10.1016/j.gene.2005.03.001;
Yang S., Liu Y., Lin A.A., Cavalli-Sforza L.L.,
"Adaptive evolution of MRGX2, a human sensory ne
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (OCT-2004) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                   Name=MRGX2;
                                                                                                                                                                                                                                                                                                                                                                                                                                             MRGX2.
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Taylor N., Ford R., i
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5 (TrEMBLrel.
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75.8%; Pred. No. 7.
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RESULT 11
Q4QXU2_PYGBI PRELIMINARY;
AC Q4QXU2;
AC Q4QXU2;
AC Q4QXU2;
AC Q4QXU2;
DT 13-SEP-2005 (TYEMBLrel. 31,
DT MRGX2;
GN MCGALT TA SET (Black snub-
CC Elwaryota; Metazoa; Chordato
CC Elwaryota; Metazoa; Chordato
CC Elwaryota; Butheria; Euarchoi)
CC Elwaryota; Metazoa; Colobinae;
CC Mammalia; Butheria; Euarchoi)
CC Cercopithecidae; Colobinae;
OX NCBI_TAXID=61621;
RN NCCLEOTIDE SEQUENCE.
RP NUCLEOTIDE SEQUENCE.
RT "Adaptive evolution of MRGX:
Involved in nociception.";
RI Gene 352C:30-35 (3005).
CC -1- SUBCELLULAR LOCATION: II
DR EMBL; AX651166; AAW70079.1;
DR PENDIT; PRO0237; GPCRRHODOP;
DR PRONSITE; PR00237; GPCRRHODOP;
DR PROSITE; PS00237; GPCRRHODOP;
DR PROSITE; PS00237; GPCRTEIN
DR PROSITE; PS00237; GPCRTEIN
DR PROSITE; PS00237; GPCRRHODOP;

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Best Local Sim
Matches 210;
    x PubMed=15662286; DOI:10.1016/j.gene.2005.03.001;
x Yang S., Liu Y., Lin A.A., Cavalli-Sforza L.L., Zhao Z.,
x Yang S., Liu Y., Lin A.A., Cavalli-Sforza L.L., Zhao Z.,
x Yang S., Liu Y., Lin A.A., Cavalli-Sforza L.L., Zhao Z.,
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x Yang S., Liu Y., Lin A.A., Cavalli-Sforza L.L., Zhao Z.,
x Yang S., Liu Y., Lin A.A., Gavalli-Sforza L.L.,
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Pfam; PF00021; 7tm 1; 1.

PRINTS; PR00237; GPCRHODOPSN.

PROSITE; PS00237; G_PROTEIN_RECEP_F1_1; UNKNOWN_1.

PROSITE; PS00237; G_PROTEIN_RECEP_F1_2; 1.

G-protein_coupled_receptor; Receptor; Transducer; TransG-protein_coupled_receptor; Transducer; TransG-protein_coupled_receptor; Transducer; T
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SUBCELLULAR LOCATION: Integral membrane
L; AY651164; AAW70077.1; -; Genomic_DNA.
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No. 1.3e-63
                                                             Transducer; Transmembrane
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Query Match

59

9

Score

998.5;

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2

Length 330;

Extracellular (Potential).

1 (Potential).

Cytoplasmic (Potential).

2 (Potential).

Transducer; Transmembrane.

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Matches 208; Conservative
EMBL; AY772458; AAV49125.1; -; Genomic_DNA
InterPro; IPR000276; GPCR_Rhodpsn.
Pfam; PF00001; 7tm 1; 1.

PRINTS; PR00237; GPCRRHODDPSN.
PROSITE; PS00237; GPCRRHODDPSN.
PROSITE; PS00237; GPROTEIN_RECEP_F1_2; 1.
G-protein coupled receptor; Receptor; Transpo_DOM 34 54 1 (Potential)
TOPO_DOM 55 63 Cytoplasmic (1)
TRANSMEM 34 84 2 (Potential)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MRGX2_MACFA
Q5U9D9;
10-MAY-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-MAY-2005 (Rel. 47, Created)
10-MAY-2005 (Rel. 47, Last sequence update)
13-SEP-2005 (Rel. 48, Last annotation update)
Mas-related G-protein coupled receptor member X2.
Name=MRGPRX2; Synonyms=MRGX2;
Macaca fascicularis (Crab eating macague) (Cynomolgus monkey).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MRGX2_N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (OCT-2004) to the EMBL/GenBank/DDBJ databases - II- FUNCTION: Orphan receptor. Probably involved in the nociceptive neurons. May regulate nociceptor function development, including the sensation or modulation of the sensation of the s
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Zhang L., Taylor N., Ford R., Conting and expression of MRG
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Mammalia; Eutheria; Euarchontoglires; Primates;
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NCBI_TaxID=9541;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   s Swiss-Prot entry is copyright. It is produced through ween the Swiss Institute of Bioinformatics and the EM European Bioinformatics Institute. There are no restr as long as its content is in no way modified and this s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SUBCELLULAR LOCATION: Integral membrane protein. SIMILARITY: Belongs to the G-protein coupled rec
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0; Mismatches
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ches 82;
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                                                                                                                                                                                                          FALSE_NEG
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Eukaryota; Metazoa; Chordata; Craniata; ru
Mammalia; Eutheria; Euarchontoglires; Prin
Mammalia; Eutheria; Cercopithecinae; Macaca.
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FROSTIE; PSO262; G_PROTEIN_RECEP_F1_2; 1.

G-protein coupled receptor; Receptor; Transducer SEQUENCE 329 AA; 37041 MM; 02AB400A7BC7EB99
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                                                                                                                                                                        involved in nociception.";
Gene 352C:30-35(2005).
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Best Local Similarity
Matches 209; Conser
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25-OCT-2004
25-OCT-2004
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25-OCT-2004 (Rel. 45, Last sequence update)
25-OCT-2005 (Rel. 48, Last annotation update)
13-SEP-2005 (Rel. 48, Last annotation update)
Mas-related G-protein coupled receptor member X2.
Name-MRGPRX2; Synonyms=MRGX2;
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Homin
MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899; Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler (Klausner R.D., Ceberg B., Buetow K.H., Schaefer C.F., Bhat N.I Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Hat N.I Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F., Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
                                                                                                                                                                                                                                                                                       Suwa M., Sato T., Okouc
Tsutsumi S., Aburatani
"Genome-wide discovery
                                                                                                                                                                                                                                                                                                                                                                                                                          genome sequence.";
FEBS Lett. 520:97-101(2002)
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MEDLINS=22040266, PubMed=12044878; DOI=10.1016
Takeda S., Kadowaki S., Haga T., Takaesu H., M
"Identification of G protein-coupled receptor
genome sequence.";
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MEDLINE=21435808; PubMed=11551509; DOI=10.1016/S0092-8674(01)00483-4; Dong X., Han S.-K., Zylka M.J., Simon M.I., Anderson D.J.; Dong X. and S.-K., Zylka M.J., Simon M.I., Enderson D.J.; "A diverse family of GPCRs expressed in specific subsets of
                                                                                                                                                                                                                                        receptor genes.";
Submitted (JUL-20
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Cell 106:619-632(2001).
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atani H., Asai K., Akiyama y
overy and analysis of human
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ki S., Haga T., Takaesu H., Mitaku S.;
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yama Y.;
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Ensembl; ENSG00000183695; Homo sapie HGNC; HGNC:17983; MRGPRX2.

MIM; 607228; -...
GO:0006021; C:integral to membra GO; GO:0004930; F:G-protein coupled GO; GO:0042923; F:neuropeptide bindi GO; GO:0042923; P:perception of pain GO; GO:0019233; P:perception of pain GO; GO:0030431; P:sleep; NAS.

InterPro; IPR000276; GFCR_Rhodpsn.
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Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.
Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
Vallalon D.K., McEwan M., Madan A., Rodrigues S., Sanchez A.
Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
"Generation and initial analysis of more than 15,000 full-length hum
"Generation and initial analysis of more than 15,000 full-length hum
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J. Biol. Chem.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and mouse cDNA sequences."; 
Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002)
                                                                                                                                                     Transmembrane.
                                                                                                                                                                                                                                   Pfam; PF00001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              removed
                                                                                                                                                                                                                                                              GO:0016021; C:integral to membrane; IC. GO:0016021; C:integral to membrane; IC. GO:0004930; F:G-protein coupled receptor GO:0042923; F:neuropeptide binding; IPI. GO:0019233; P:perception of pain; NAS. GO:0030431; P:sleep; NAS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  coupled.
SUBCELLULAR LOCATION: Integral membrane protein.
TISSUE SPECIFICITY: Has a limited expression profile, both
peripheral and within the central nervous system, with highest
levels in dorsal root ganglion.
SIMILARITY: Belongs to the G-protein coupled receptor 1 family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FUNCTION: Orphan receptor. Probably involved in the function nociceptive neurons. May regulate nociceptor function and/or development, including the sensation or modulation of pain. Cortistatin-14 seems to be a high potency ligand at this reconstitution has several biological functions including roles.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        s Swiss-Prot entry is copyright. It is produced through a collaboration ween the Swiss Institute of Bioinformatics and the EMBL outstation European Bioinformatics Institute. There are no restrictions on it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sleep regulation locomotor activity, and cortical function. In receptor-expressing cells, cortistatin-stimulated increases in intracellular Ca(2+) but had no effect on basal or forskolinstimulated cAMP levels, suggesting that this receptor is G(q)-
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Toshiyuki S., Carninci P.,
Peters G.J., Abramson R.D.,
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        1 (Potential).

Cytoplasmic (Potential).
2 (Potential).
Extracellular (Potential).
3 (Potential).
Cytoplasmic (Potential).
4 (Potential).
Extracellular (Potential).
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Mullahy S.J.,
naratne P.H.,
J., Hulyk S.W.,
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Best Local &
Matches 205
involved in nociception.";

Gene 352:30-35(2005).

-!- SUBCELLULAR LOCATION: Integral membrane pr
-!- SUBCELLULAR LOCATION: Integral membrane pr
-:- SUBCELLULAR LOCATION: Integral membrane pr
-:- SUBCELLULAR LOCATION: Integral membrane pr
-:- EMBL; AY651161; AAW70073.1; -; Genomic_DNA.
-- EMBL; AY651143; AAW70056.1; -; Genomic_DNA.
-- EMBL; AY651144; AAW70057.1; -; Genomic_DNA.
-- EMBL; AY651144; AAW70082.1; -; Genomic_DNA.
-- EMBL; AY651145; AAW70082.1; -; Genomic_DNA.
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Q4QXW4;
13-SEP-2005 (TrEMBLrel. 3:
13-SEP-2005 (TrEMBLrel. 3:
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PubMed=15862286; DOI=10.1016/j.gene.2005.03.001;

PubMed=15862286; DOI=10.1016/j.gene.2005.03.001;

PubMed=15862286; DOI=10.1016/j.gene.2005.03.001;

PubMed=15862286; DOI=10.1016/j.gene.2005.03.001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens (Human).
Eukaryota; Metazoa; (
Mammalia; Eutheria; E
                                                                                                                                                                                         Yang S., Liu Y., Lin A.A., Ca
"Adaptive evolution of MRGX2,
                                                                                                                                                                                                                                     NUCLEOTIDE
                                                                                                                                                                                                                                                                            involved in nociception.";
Gene 352C:30-35(2005).
                                                                                                                                                                                                                                                                                                                        Yang S., Liu Y., Lin A.A., Cavalli-Sforza L.L., Zhao
"Adaptive evolution of MRGX2, a human sensory neuron
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                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=9606;
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IAMLVFLCVVLCGSSLVLLVRILCGSRKMPLTRLYVTILLTVLVFLLCGLPFGIQWALFS 	CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRGILEWMPCDFLE	IYILNLVAANFLFLSGHIIFSPLPLINIRHPISKILSPVWTF	ETPCYNQTLSFTGLTCIIS	%; Score 1691; DB 2; Le %; Pred. No. 8.4e-138; 0; Mismatches 0; In	ein-Coupled Receptors from 17 US/09/254,227A 3-03	254227A	ALIGNMENTS	US-08-417-103-6 US-08-417-103-16 US-09-170-496D-164	US-09-576-160B-6 US-07-816-283-6	US-09-456-455A-3 US-08-676-351-5	US-09-745-842-21 US-09-261-599B-3	US-09-826-509-501 US-09-170-496D-2	US-09-944-807-2	US-08-202-056-7	525-573-8	-390-000A -446-669-
VFLLCGLPFGIQWALFS 240	LFSGANSVWCETSDFIT 180	FPYFIGLSMLSAISTER 120	LVALTGNAVVLWLLGCRMRRNA 60	Length 322; Indels 0; Gaps 0	the Rat and Human			Sequence 6, Appli Sequence 16, Appl Sequence 164, Appl	<i>ي</i> ف	'n'n	ω <sub>21</sub>	2 50	ν. <sub>+</sub>		φ.	nce 5,

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APPLICANT: Port, Krishna G.
APPLICANT: Moffett, Serge
APPLICANT: Moffett, Serge
APPLICANT: Abran, Daniel
TITLE OF INVENTION: METHODS AND COMPOUNDS FOR PREVENTION AND TREATMENT OF ELEVATED
TITLE OF INVENTION: INTRACCULAR PRESSURE AND RELATED CONDITIONS
FILE REFERENCE: 4518/1M674US1
FILE REFERENCE: 4518/1M674US1
CURRENT APPLICATION NUMBER: US/10/401,397A
CURRENT FILING DATE: 2003-03-27
PRIOR FILING DATE: 2002-03-27
PRIOR FILING DATE: 2002-03-27
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin version 3.1
SOFTWARE: E
SEQ ID NO 5
LENGTH: 322
                                           GENERAL INFORMATION:

APPLICANT: Ahmad, Sultan
APPLICANT: Banville, Denis
APPLICANT: Banville, Denis
APPLICANT: Benotin, Yves
APPLICANT: Lembo, Paola
APPLICANT: O'Donnell, Dajan
APPLICANT: O'Donnell, Dajan
APPLICANT: Shi-Hsiang, Shen
TITLE OF INVENTION: G Protein-Coupled Receptors from the Rat and Human
FILE REFERENCE: 81823/268117
CURRENT APPLICATION NUMBER: US/09/254,227A
CURRENT FILING DATE: 1999-03-03
NUMBER OF SEQ ID NOS: 22
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LENGTH: 322
TYPE: PRT
ORGANISM: Homo sapiens
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Patent No. 6864239
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Best Local Similarity 96.9%;
Matches 312; Conservative
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                                     PatentIn version 3.0
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Pred. No. 3.7e-133;
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; SEQ ID NO 7
; LENGTH: 322
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-254-227A-7
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; ORGANISM: Homo sapiens
US-09-254-227A-5
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Best Local Similarity 94.1%;
Matches 302; Conservative
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Best Local
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APPLICANT: Lembo, Paola
APPLICANT: O'Donnell, Dajan
APPLICANT: Shi-Hslang, Shen
TITLE OF INVENTION: G Protein-Coupled Receptors from
FILE REFERENCE: 81823/268117
CURRENT APPLICATION NUMBER: US/09/254,227A
CURRENT FILING DATE: 1999-03-03
NUMBER OF SEQ ID NOS: 22
NUMBER OF SEQ ID NOS: 22
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                                             121 CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSDFIT 180
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                                                                                          61 FSIYILNLAAADFLFLSGRLIYSLLSFISIEHTISKILYFVVMFSYFAGLNFLSAVSTDR 120
                                                                                                             61 VSIYILNLVAANFLFLSGHIIFSPLFLINIRHPISKILSPVMTFPYFIGLSMLSAISTER
                                                                                                                                                       Similarity
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IAWLVFLCVVLCGSSLVLLVRILCGSRKMPLTRLYVTILLTVLVFLLCGLPFGIQWALFS
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82.3%; Pred. No. 2.5e-112;
ative 24; Mismatches 33;
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Pred. No. 1.8e-127;
8; Mismatches 11;
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SEQ ID NO 9
LENGTH: 322
TYPE: PRT
ORGANISM: Homo sapiens
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Sequence 20, Application US/10314048A
Patent No. 6902902
GENERAL INFORMATION:
APPLICANT: Unett, David J.
APPLICANT: Chen, Ruoping
APPLICANT: Richman, Jeremy
APPLICANT: Connolly, Daniel
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GENERAL INFORMATION:
APPLICANT: Ahmad, Sultan
APPLICANT: Banville, Denis
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APPLICANT: Lembo, Paola
APPLICANT: Lembo, Paola
APPLICANT: O'Donnell, Dajan
APPLICANT: Shi-Hsiang, Shen
TITLE OF INVENTION: G Protein-Coupled Receptors from PILE REFERENCE: 81823/268117
CURRENT APPLICATION NUMBER: US/09/254,227A
CURRENT FILING DATE: 1999-03-03
NUMBER OF SEQ ID NOS: 22
SOFTWARE: Patentin version 3.0
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82.3%; Pred. No. 4e-111;
tive 22; Mismatches 3
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APPLICANT: Liaw, Cuest
APPLICANT: Lowitz, Revin P.
APPLICANT: Behan, Dominic P.
APPLICANT: Behan, Dominic P.
APPLICANT: Chalmers, Derek T.
APPLICANT: Lerner, Michael
TITLE OF INVENTION: Human G Protein-Coupled Receptors and Modulators Thereof
TITLE OF INVENTION: for the Treatment of Metabolic-Related Disorders
PILE REFERENCE: 22.US6.CIP
CURRENT APPLICATION NUMBER: US/10/314,048A
CURRENT PILING DATE: 2002-12-06
PRIOR APPLICATION NUMBER: 10/096.511
PRIOR FILING DATE: 2002-03-12
PRIOR APPLICATION NUMBER: 09/995,543
PRIOR APPLICATION NUMBER: 09/995,543
PRIOR APPLICATION NUMBER: 60/399,917
PRIOR APPLICATION NUMBER: 60/399,917
PRIOR APPLICATION NUMBER: 60/399,917
PRIOR APPLICATION NUMBER: 60/399,917
PRIOR APPLICATION NUMBER: 60/404,761
PRIOR FILING DATE: 2002-07-29
PRIOR APPLICATION NUMBER: 60/404,761
PRIOR FILING DATE: 2002-09-13
NUMBER OF SEQ ID NOS: 161
SOFTWARE: Patentin version 3.1
SOFTWARE: Patentin version 3.1
SOFTWARE: Patentin version 3.1
SOFTWARE: Patentin version 3.1
SOFTWARE: OSC 10 NO 20
LENGTH: 322
TYPE: PRT
ORGANISM: Homo sapiens
US-10-314-048A-20
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US-09-254-227A-13
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Sequence 13, Application US/09254227A
Patent No. 6659257
GENERAL INFORMATION:
APPLICANT: Abmad, Sultan
APPLICANT: Banville, Denis
APPLICANT: Fortin, Yves
APPLICANT: Lembo, Paola
APPLICANT: O'Donnell, Dajan
APPLICANT: Shi-Hsiang, Shen
TITLE OF INVENTION: G Protein-Coupled Receptors from the
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Best Local
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Liaw, Chen
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rd, James
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82.0%; Pred. No. 1.9e-110;
vative 22; Mismatches 36;
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CURRENT APPLICATION NUMBER: US/09/254,227A; CURRENT FILING DATE: 1999-03-03; NUMBER OF SEQ ID NOS: 22; SOFTWARE: Patentin version 3.0; SEQ ID NO 13; LENGTH: 322; TYPE: PRT; ORGANISM: Homo sapiens
US-09-254-227A-13
                                                                                                                                                                                    ; SOFTWARE: PatentIn vers
; SEQ ID NO 11
; LENGTH: 322
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-254-227A-11
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US-09-254-227A-11
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Best Local Simi
Matches 264;
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APPLICANT: Banville, Denis
APPLICANT: Fortin, Yves
APPLICANT: Fortin, Yves
APPLICANT: O'Donnell, Dajan
APPLICANT: O'Donnell, Dajan
APPLICANT: Shi-Hsiang, Shen
TITLE OF INVENTION: G Protein-Coupled Receptors from the Rat and Human
FILE REFERENCE: 81823/268117
CURRENT APPLICATION NUMBER: US/09/254,227A
CURRENT FILING DATE: 1999-03-03
NUMBER OF SEQ ID NOS: 22
NUMBER OF SEQ ID NOS: 22
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Local Similarity 83.1%;
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Similarity 82.5%;
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MDPTIPVLGTKLTPINGREETPCYNQTLSFTGLTCIISLVALTGNAVVLWLLGCRMRRNA
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                                                                                                                   ; Score 1356; DB 2;
; Pred. No. 5.6e-109;
21; Mismatches 35;
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Pred. No. 9.4e-110;
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CURRENT APPLICATION NUMBER: US/10/314,048A
CURRENT FILING DATE: 2002-12-06
PRIOR APPLICATION NUMBER: 10/096,511
PRIOR FILING DATE: 2002-03-12
PRIOR APPLICATION NUMBER: 09/995,543
PRIOR PILING DATE: 2001-11-27
PRIOR APPLICATION NUMBER: 09/39,917
PRIOR PILING DATE: 2002-07-29
PRIOR PILING DATE: 2002-07-29
PRIOR PILING DATE: 2002-08-19
PRIOR PILING DATE: 2002-08-19
PRIOR PILING DATE: 2002-09-13
PRIOR FILING DATE: 2002-09-13
NUMBER OF SEQ ID NOS: 161
SOPTWARE: PATENTIN VERBION 3.1
SEQ ID NO 30
LENGTH: 330
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US-10-314-048A-30
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US-10-314-048A-30
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                                                                                                                                                                                                                                                                                                                                           Query Match 57.9%; Score 978.5; DB 2; Best Local Similarity 62.3%; Pred. No. 1.7e-76; Matches 205; Conservative 29; Mismatches 86;
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APPLICANT: Behan, Dominic P.
APPLICANT: Chalmers, Derek T.
APPLICANT: Chalmers, Michael
APPLICANT: Lerner, Michael
APPLICANT: Lerner, Michael
APPLICANT: Lerner, Michael
APPLICANT: MICHAEL
APPLICANT: Chalmers, Michael
APPLICANT: Chalmers, Michael
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APPLICANT: Chalmers, Michael
APPLICANT: Levis, 
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APPLICANT:
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APPLICANT: Chen, Ruoping
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114 SAISTERCLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWC 173
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                                                                    61 RNAFSVYVLSLAGADFLFLCFQIINCLVYLSNFFCSISINFPSFFTTVMTCAYLAGLSML
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                                                                                                    RNAVSIYILNLVAANFLELSGHIIFSPLFLINIRHFIS----KILSPVMTFFYFIGLSML
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                                                                                                                                                                                                             MDPTTPAWGTESTTVNGNDQALLLLCGKETLIPVFLILFIALVGLVGNGFVLMLLGFRMR
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Leonard, James
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Liaw, Chen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Connolly, Daniel
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APPLICANT: Ahmad, Sultan
APPLICANT: Banville, Denis
APPLICANT: Fortin, Yves
APPLICANT: Lembo, Paola
APPLICANT: Lembo, Paola
APPLICANT: O'Donnell, Dajan
APPLICANT: Shi-Hsiang, Shen
TITLE OF INVENTION: G Protein-Coupled Receptors fro
FILE REFERENCE: 81823/268117
CURRENT APPLICATION NUMBER: US/09/254,227A
CURRENT FILING DATE: 1999-03-03
NUMBER OF SEQ ID NOS: 22
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1
LENGTH: 337
TYPE: PRT
ORGANISM: rat
          RESULT 11
US-10-314-048A-10
; Sequence 10, Application US/10314048A
; Patent No. 6902902
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US-09-254-227A-1
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Patent No. 6696257
GENERAL INFORMATION:
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 48.2%; Score 815.5; DB 2; Best Local Similarity 53.4%; Pred. No. 1.9e-62;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches 173;
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                                                                                                                                       RALQDTPEVDEGGGWLPQETLELS 316
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                                                                                                                    RALEETPEEDEYTDSHVQKPTEIS 333
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;Patent No.
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PRIOR APPLICATION NUMBER: 10/096,511
PRIOR FILING DATE: 2002-03-12
PRIOR PLILING DATE: 2002-01-12
PRIOR PLILING DATE: 2001-11-27
PRIOR APPLICATION NUMBER: 60/399,917
PRIOR FILING DATE: 2002-07-29
PRIOR APPLICATION NUMBER: 60/404,761
PRIOR APPLICATION NUMBER: 60/404,761
PRIOR APPLICATION NUMBER: 60/404,761
PRIOR APPLICATION NUMBER: 60/410,747
PRIOR APPLICATION NUMBER: 60/410,747
PRIOR PRILING DATE: 2002-08-19
PRIOR PRILING DATE: 2002-09-13
PRIOR FILING DATE: 2002-09-13
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              OLTAVIO

TITLE OF INVENTION: DNA SEQUENCES ENCODING MAS ONHCOGENE,
POLYPETIDES ENCODED THEREFROM AND DIAGNOSTIC AND OTHER METHODS
BASED THEREFROM
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Best Local Simi
Matches 127;
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SOFTWARE: Patentin version 3.1
SEQ ID NO 10
LENGTH: 321
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APPLICANT: Chen, R
APPLICANT: Richman
APPLICANT: Connoll
APPLICANT: Dang, H
APPLICANT: Choi, B
APPLICANT: Leonard
APPLICANT: Hakak,
APPLICANT: Liaw, C
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APPLICANT:
APPLICANT:
APPLICANT:
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ORGANISM: Homo sapiens
                                                                APPLICANT: Young, Dallan; Wigler, Michael H.; Fasano
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                                                                                                                                                                                                  291 LORALODTPEVDEGG 305
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Choi, Bryan
Leonard, James
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Liaw, Chen
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Connolly, Daniel
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APPLICANT: Chalmers, Derek T.
APPLICANT: Lerner, Michael
TITLE OF INVENTION: Human G Protein-Coupled Receptors and Modulators Thereof
TITLE OF INVENTION: for the Treatment of Metabolic-Related Disorders
FILE REFERENCE: 22.US6.CIP
180 MVQAALIMGVLTPVMTLSSLTLEVMVRRSSQQWRRQP-TRLEVVVLASVLVELICSLPLS 238
                                                                                                                         178 FITIAMLY-FLCVVLCGSSLVLLVRILCGS---RKMPLTRLYVTILLTVLVFLLCGLPFG
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                                                                                                                                                                                                                            118 TERCLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSD
                                                                                                                                                                                                                                                                                     61 FCIYILMLAAADLLFLFSMASTLSLETQPLVNTTDKVHELMKRLMYFAYTVGLSLLTAIS
                                                                                                                                                                                                                                                                                                                                      61 VSIYILNLVAANFLFL---SGHIIFSPLPLINIRHPISKILSPVMTFPYFIGLSMLSAIS 117
                                                                                                                                                                                                                                                                                                                                                                                                                          1 MDFTIPVLGTKLTPINGREETPCYNQTLSFTGLTCIISLVALTGNAVVLWLLGCRMRRNA
                                                                                                                                                                                                                                                                                                                                                                                NNQTLNSSGTVESALNYSRGSTVHTAYLVLSSLAMFTCLCGMAGNSMVIWLLGFRWHRNP
                                                                                                                                                                                         30.2%; Score 510; DB 2; Length 321;
40.3%; Pred. No. 3.4e-36;
vative 57; Mismatches 115; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16;
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NUMBER OF SEQUENCES: 2
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/
FILING DATE: 06-JUN-1986
SEQ ID NO:2:
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US-08-118-270-52
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Best Local Similarity
Matches 105; Conserv
                                                                    COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/118,270
FILING DATE: 09-SEP-193
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/943,236
FILING DATE: 10-SEP-1992
ATTORNEY, AGENT INFORMATION:
NAME: TOWNSEND, NUMBER: MURPHY=2A
REFERENCE/DOCKET NUMBER: MURPHY=2A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No. 5508384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Schuster, David I.
TITLE OF INVENTION: POLYPEPTIDES OF G-COUPLED PROTEIN
TITLE OF INVENTION: RECEPTORS, AND COMPOSITIONS AND METHODS THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Murphy, Randall B. APPLICANT: Schuster, David I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                  TELECOMMUNICATION INFORMATION: TELEPHONE: 202-628-5197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 325
TELEPHONE: 202-01.
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ZIP: 20004
                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 419 Seven
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               298
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TVYITHLSIADISLLFCIFILSIDYALDYELSSGHYYTIVTLSVTFLFGYNTGLYLLTAI 126
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419 Seventh Street, N.W., Suite 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24.6%; Score 415.5; DB 6; nilarity 34.1%; Pred. No. 4.7e-28; Conservative 64; Mismatches 112;
                                                                                                                                                                                                                                                                                                                                                                                           USA
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SEQUENCE CHARACTERISTICS
                        TELEX: 248633
INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
LENGTH: 282 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: New York University
TITLE OF INVENTION: POLYEPTIDES OF G-COUPLED PROTEIN
TITLE OF INVENTION: RECEPTORS, AND COMPOSITIONS AND METHODS THEREOF
NUMBER OF SEQUENCES: 348
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE:
                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/943,236
FILING DATE: 10-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Townsend, Kevin G.
REGISTRATION NUMBER: 34,033
REFERENCE/DOCKET NUMBER: MURPHY=2 PCT
                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARB: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
TELEFAX: 202-737-3528
                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Washington
STATE: D.C.
COUNTRY: USA
ZIP: 20004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 282 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      / Match 22.9%; Score 386.5; DB 1; Local Similarity 34.4%; Pred. No. 1.3e-25;
               TYPE: amino acid
                                                                                                                                                                                                                                                                            FILING DATE: 09-SEP-1993
STRANDEDNESS:
                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             168
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          224 LFSTINSSANPFIYFFVGSSKKKRFKESLKVVLTRAFKD 262
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    96;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             VVKIRKNTWASHSSKLYIVIMVTIIIFLIFAMPMRLLYLLYYEY---WST-FGNLHHISL
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419 Seventh Street, N.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Suite 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         96;
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RESULT 15
US-08-118-270-76
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 76, Application US/08118270
Patent No. 5508384
GENERAL INFORMATION:
APPLICANT: Murphy, Randall B.
APPLICANT: Schuster, David I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 34.4
Matches 96; Conservative
                                                                                                                                                                                                                                                                      COMPUTER: IBM PC COMPARTIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/118,270
FILING DATE: 09-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/943,236
FILING DATE: 10-SEP-1992
ATTORNEY/AGENT INFORMATION:
MOLECULE TYPE: peptide -08-118-270-76
                                                                                   TELEFAX: 202-737-3528
TELEX: 248633
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 298 amino acid
                                                                                                                                                                                   TELECOMMUNICATION INFORMATION: TELEPHONE: 202-628-5197
                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy COMPUTER: IBM PC com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Murphy, Randall B.

APPLICANT: Schuster, David I.

TITLE OF INVENTION: RECEPTORS, AND COMPOSITIONS AND METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                  TELEPHONE: 202-0-3528
                                                                                                                                                                                                                    NAME: Townsend, Kevin G.
REGISTRATION NUMBER: 34,033
REFERENCE/DOCKET NUMBER: MURPHY=2A
                                                  TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 419 Seven
                                   TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   224 LFSTINSSANPFIYFFVGSSKKKRFKESLKVVLTRAFKD 262
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         168 VVKIRKNTWASHSSKLYIVIMVTIIIFLIFAMPMRLLYLLYYEY---WST-FGNLHHISL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    139 SSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSDFITIAWLVFLCVVLCGSSLVL 198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    259 FLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQD 297
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         37 ISLVALTGNAVVLW--------LIGCRMRRNAVSIYILNL-VAANFLFLSG
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Best Local Similarity
Matches 91; Conserv
234 CICINSSAKPIVYFIAGRDKSQRLMEPLRVVFQRALRDGAEPGDAASSTENTVIMEM 290
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                                           260 LSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPEVDEGGGWLPQE-TLEL 315
                                                                                    180 ECRARRQRSAKLNHVVLAIVSVFLVSSIYLGIDWFLF-----WVFQIPAPFPEYVRDL
                                                                                                                           203 LCGSRKMPLT-RLYVTILLTVLVFLLCGLPFGIQWALFSRIHLDW--KVLFCHVHLVSIF 259
                                                                                                                                                                                                             148 ALSLLRSILEWMFCDFLFSGANSVWCETSDFITIAWLVF-----LCVVLCGSSLVLLVRI 202
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Search completed: February 3, 2006, 20:33:05
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/cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*
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US-11-083-611-31
US-11-083-611-31
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US-10-957-135-31
US-10-957-135-31
US-10-977-810-2
US-10-977-810-2
US-10-973-810-2
US-10-213-84-7-9
US-10-231-074-2
US-10-231-074-2
US-10-231-074-2
US-10-237-467-10
US-10-183-116-16
US-10-183-116-16
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ς.	Sequence 33, Appl	Sequence 18, Appl	Sequence 18, Appl	Sequence 532, App	Sequence 528, App	Sequence 4, Appli	Sequence 16, Appl	Sequence 20, Appl		Sequence 20, Appl	Sequence 2, Appli	Sequence 20, Appl	Sequence 20, Appl	Sequence 44, Appl	•	Sequence 527, App	-

## ALIGNMENTS

RESULT 1 US-10-292-798-1274

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CURRENT APPLICATION NUMBER: US/10/292,798
CURRENT FILING DATE: 2002-11-13
PRIOR APPLICATION NUMBER: 10/017,161
PRIOR FILING DATE: 2001-12-18
PRIOR APPLICATION NUMBER: JP 2001-246789
PRIOR PILING DATE: 2001-06-18
PRIOR FILING DATE: 2001-05-18
NUMBER OF SEQ ID NOS: 2070
SOFTWARE: PAtentin Ver. 2.1
SEQ ID NO 1274
LENGTH: 322
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Best Local Similarity
Matches 314; Conserv
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APPLICANT: ASAI, KIYOSHI
APPLICANT: AKIYAMA, YUTAKA
APPLICANT: ABURATANI, HIROYUKI
TITLE OF INVENTION: GUANOSINE TRIPHOSPHATE-BINDING PROTEIN COUPLED RECEPTORS
FILE REFERENCE: 084335/166
                                                                      241
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                                                     RIHLDWKVLFCHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPE
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                                  RIHLDWKVLFCHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPE
                                                                                                          IAWLVFLCVVLCGSSLVLLVRILCGSRKMPLTRLYVTILLTVLVFLLCGLPFGIQWALFS
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; Remaining Prior Application data removed -
; NUMBER OF SEQ ID NOS: 1391
; SEGTWARE: PatentIn Ver. 2.1
; SEQ ID NO 530
; LENGTH: 322
; TYPE: PRT
; ORCANISM: Homo sapiens
US-10-072-012-530
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APPLICANT: Burgess, Catherine E.

APPLICANT: Burgess, Catherine E.

TITLE OF INVENTION: Proteins and Nucleic Aci

FILE REFERENCE: 21402-258

CURRENT APPLICATION NUMBER: US/10/072,012

CURRENT FILING DATE: 2002-01-31

PRIOR PELLING DATE: 2001-01-30

PRIOR FILING DATE: 2001-01-31

PRIOR FILING DATE: 2001-01-31

PRIOR FILING DATE: 2001-01-31

PRIOR APPLICATION NUMBER: 60/265,517

PRIOR FILING DATE: 2001-01-31

PRIOR APPLICATION NUMBER: 60/265,412

PRIOR APPLICATION NUMBER: 60/265,412

PRIOR APPLICATION NUMBER: 60/265,395

PRIOR APPLICATION NUMBER: 60/265,395

PRIOR APPLICATION NUMBER: 60/265,395

PRIOR FILING DATE: 2001-01-31

PRIOR APPLICATION NUMBER: 60/265,395

PRIOR FILING DATE: 2001-01-31

PRIOR APPLICATION NUMBER: 60/265,395

PRIOR FILING DATE: 2001-01-31

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PRIOR APPLICATION NUMBER: 60/265,412
PRIOR TILING DATE: 2001-01-31
PRIOR PELICATION NUMBER: 60/265,395
PRIOR PELICATION NUMBER: 60/266,395
PRIOR PELICATION NUMBER: 60/266,406
PRIOR APPLICATION NUMBER: 60/266,767
PRIOR PILING DATE: 2001-02-02
PRIOR PILING DATE: 2001-02-07
PRIOR PELICATION NUMBER: 60/267,057
PRIOR APPLICATION NUMBER: 60/266,975
PRIOR PELICATION NUMBER: 60/266,975
PRIOR APPLICATION NUMBER: 60/267,459
PRIOR APPLICATION NUMBER: 60/267,459
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PRIOR APPLICATION NUMBER: 60/267,459
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                                                                                        Local Similarity
nes 314; Conserv
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INVENTION: Proteins and Nucleic Acids Encoding
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Patturajan, Muu-
Patturajan, Muu-
Patta, Richard
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Taupier Jr, Raymond
Gusev, Vladimir Y.
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Rieger, Daniel K.
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Grosse, William M.
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Zerhusen, Bryan
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                                                                                        97.4%;
llarity 97.5%;
Conservative
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David W.
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                                                                                        Score 1647; DB 4;
Pred. No. 8.9e-140;
4; Mismatches 4;
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                                                                                                                                                                                                                                                                                                                                          See File Wrapper or PALM
                                                                                                                                  Length 322;
                                                                                          Indels
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FILE REFERENCE: 21402-258
CURRENT APPLICATION NUMBER: US/10/072,012
CURRENT FILING DATE: 2002-01-31
PRIOR APPLICATION NUMBER: 60/265,102
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: 60/265,514
PRIOR APPLICATION NUMBER: 60/265,517
PRIOR APPLICATION NUMBER: 60/265,517
PRIOR APPLICATION NUMBER: 60/265,412
PRIOR FILING DATE: 2001-01-31
PRIOR APPLICATION NUMBER: 60/265,412
PRIOR APPLICATION NUMBER: 60/265,412
PRIOR APPLICATION NUMBER: 60/265,395
PRIOR FILING DATE: 2001-01-31
PRIOR PPLICATION NUMBER: 60/266,767
PRIOR FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: 60/266,767
PRIOR APPLICATION NUMBER: 60/266,767
PRIOR APPLICATION NUMBER: 60/266,767
PRIOR APPLICATION NUMBER: 60/266,767
PRIOR PRILING DATE: 2001-02-07
PRIOR PRILING DATE: 2001-02-07
PRIOR FILING DATE: 2001-02-07
PRIOR FILING DATE: 2001-02-07
PRIOR PRILING DATE: 2001-02-07
PRIOR APPLICATION NUMBER: 60/266,975
PRIOR APPLICATION NUMBER: 60/266,975
PRIOR APPLICATION NUMBER: 60/267,459
PRIOR APPLICATION NUMBER: 60/267,459
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APPLICANT: Tchernev,
APPLICANT: Spytek,
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APPLICANT:
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APPLICANT: Rieger, Daniel K.
APPLICANT: Burgess, Catherine E.
TITLE OF INVENTION: Proteins and Nucleic Acids Encoding
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Gerlach, Valerie
Taupier Jr, Raymond J.
Gusev, Vladimir Y.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Furtak, Katarzyna
Grosse, William M.
Alsobrook II, John P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pena, Carol E. A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wolenc, Adam R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Padigaru, Muralidhara
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TITLE OF INVENTION: Receptors
FILE REFERENCE: AREN-0308
FILE REFERENCE: AREN-0308
CURRENT APPLICATION NUMBER: US/09/95,225
CURRENT APPLICATION NUMBER: US/09/95,225
CURRENT FILING DATE: 2001-11-26
PRIOR APPLICATION NUMBER: 09/170,496
PRIOR FILING DATE: 1998-10-13
PRIOR APPLICATION NUMBER: PCT/US99/23938
PRIOR APPLICATION NUMBER: 60/253,404
PRIOR FILING DATE: 1998-10-13
PRIOR APPLICATION NUMBER: 60/253,404
PRIOR FILING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: 60/255,366
PRIOR FILING DATE: 2000-12-12
PRIOR APPLICATION NUMBER: 60/25,366
PRIOR APPLICATION NUMBER: 60/270,286
PRIOR APPLICATION NUMBER: 60/270,286
PRIOR APPLICATION NUMBER: 60/282,365
PRIOR APPLICATION NUMBER: 60/282,365
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NUMBER OF SEQ ID NOS: 1391
SOFTWARE: PatentIn Ver. 2:1
SEQ ID NO 535
LENGTH: 322
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 97.4%;
Best Local Similarity 97.5%;
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Chu, Zhi Liang
Dang, Huong T.
Lowitz, Kevin P.
Pride, Cameron
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSDFIT
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Pred. No. 8.9e-140;
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APPLICANT: LOWITZ, KEVIN P.

APPLICANT: Pride, Cameron

ITITLE OF INVENTION: Endogenous And No. US20

ITITLE OF INVENTION: Receptors

FILE REFERENCE: AREN-0308

CURRENT APPLICATION NUMBER: US/09/995,225

CURRENT FILING DATE: 2001-11-26

PRIOR APPLICATION NUMBER: 09/170,496

PRIOR APPLICATION NUMBER: PCT/US99/23938

PRIOR APPLICATION NUMBER: PCT/US99/23938

PRIOR FILING DATE: 1998-10-13

PRIOR FILING DATE: 1908-0-13

PRIOR FILING DATE: 2000-12-12

PRIOR APPLICATION NUMBER: 60/253,404

PRIOR FILING DATE: 2000-12-12

PRIOR APPLICATION NUMBER: 60/255,366

PRIOR APPLICATION NUMBER: 60/270,286

PRIOR APPLICATION NUMBER: 60/270,286

PRIOR APPLICATION NUMBER: 60/282,365

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PRIOR APPLICATION NUMBER: 60/282,365
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Best Local S
Matches 313
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/290,917
PRIOR FILING DATE: 2001-05-14
PRIOR APPLICATION NUMBER: 60/309,208
PRIOR FILING DATE: 2001-07-31
NUMBER OF SEQ ID NOS: 67
SOFTWARE: PatentIn version 3.1
SEQ ID NO 20
                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Chen, Ruoping
APPLICANT: Chu, Zhi Liang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE: OTHER INFORMATION: No. US20020193584A1el Sequence
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APPLICATION NUMBER: 60/270,266 FILING DATE: 2001-02-20
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vo. US20030139588A9
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Pred. No. 2.5e-139;
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Sequence 31, Application US/10183116

Publication No. US20030092035A1

GENERAL INFORMATION:

APPLICANT: Anderson, David J.

APPLICANT: Dong, Xinzhong

APPLICANT: Simon, Melvin

APPLICANT: Simon, Melvin

APPLICANT: Simon, Melvin

CURRENT APPLICATION: PAIN SIGNALING MOLECULES

FILLE REFERENCE: CALTE. 47C1CP1

CURRENT APPLICATION NUMBER: US/10/183,116

CURRENT FILING DATE: 2000-06-26

PRIOR APPLICATION NUMBER: US 60/222,344

PRIOR FILING DATE: 2000-05-04

PRIOR APPLICATION NUMBER: US 99/704,707

PRIOR APPLICATION NUMBER: US 09/704,707

PRIOR APPLICATION NUMBER: US 09/704,707

PRIOR FILING DATE: 2000-11-03

PRIOR FILING DATE: 2001-01-03

PRIOR APPLICATION NUMBER: US 09/849,869

PRIOR APPLICATION NUMBER: US 09/849,869

PRIOR APPLICATION DATE: 2001-05-04

NUMBER OF SEQ ID NOS: 109
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PRIOR APPLICATION NUMBER: 60/282,358
PRIOR APPLICATION NUMBER: 60/282,356
PRIOR APPLICATION NUMBER: 60/282,356
PRIOR PILING DATE: 2001-04-06
PRIOR PILING DATE: 2001-05-14
PRIOR PILING DATE: 2001-05-14
PRIOR APPLICATION NUMBER: 60/309,208
PRIOR APPLICATION NUMBER: 60/309,208
PRIOR FILING DATE: 2001-07-31
NUMBER OF SEQ ID NOS: 67
SOFTWARE: PATENTE APPLICATION VERSION 3.1
SEQ ID NO 20
LENGTH: 322
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Pred. No. 2.5e-139;
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US-10-225-567A-674
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APPLICANT: Burmer, Glenna C.
APPLICANT: Roush, Christine L.
APPLICANT: ROUSH, Christine L.
TITLE OF INVENTION: ANTIGENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTORS
FILE REFERENCE: 1920-4-4
FILE REFERENCE: 1920-4-4
CURRENT FILING DATE: 2001-12-19
CURRENT FILING DATE: 2001-12-19
PRIOR APPLICATION NUMBER: 60/257,144
PRIOR FILING DATE: 2000-12-19
PRIOR FILING DATE: 2000-12-19
NUMBER: OF SEQ ID NOS: 2292
SOOTWARE: Patentin version 3.1
SEQ ID NO 674
LENGTH: 322
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Best Local S
Matches 313
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SEQ ID NO 31
LENGTH: 322
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Publication No. US20
GENERAL INFORMATION:
                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                Matches 313;
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121 CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSDPIT
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                                                                 61 VSIYILNLVAANFLFLSGHIIFSFLFLINIRHFISKILSFVMTFFYFIGLSMLSAISTER
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                                                  VSIYILNLVAADFLFLSGHIICSPLRLINIRHPISKILSPVMTFPYFIGLSMLSAISTER
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o. US20030113798A1
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                                                                                                                                                                                            97.1%;
97.2%;
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                                                                                                                                                                                Score 1642; DB 4;
Pred. No. 2.5e-139;
5; Mismatches 4;
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Pred. No. 2.5e-139;
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US-10-072-012-529

GENERAL INFORMATION: Sequence 529, Appropriate Publication No.

APPLICANT:

Patturajan,

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APPLICANY: Kleger, Mainer N.

APPLICANY: Norgens, Catherine E.

APPLICANT: Burgess, Catherine E.

TITLE OF INVENTION: Proteins and Nucleic Acids Encoding Same Fills Reference: 21402-258

CURRENT ENCLOATION NUMBER: US/10/072,012

CURRENT FILING DATE: 2002-01-31

PRIOR APPLICATION NUMBER: 60/265,514

PRIOR FILING DATE: 2001-01-31

PRIOR APPLICATION NUMBER: 60/265,517

PRIOR APPLICATION NUMBER: 60/265,517

PRIOR APPLICATION NUMBER: 60/265,412

PRIOR APPLICATION NUMBER: 60/265,395

PRIOR APPLICATION NUMBER: 60/266,395

PRIOR APPLICATION NUMBER: 60/266,395

PRIOR APPLICATION NUMBER: 60/266,767

PRIOR PILING DATE: 2001-01-31

PRIOR APPLICATION NUMBER: 60/266,767

PRIOR PILING DATE: 2001-02-02

PRIOR PILING DATE: 2001-02-05

PRIOR APPLICATION NUMBER: 60/266,767

PRIOR APPLICATION NUMBER: 60/266,767

PRIOR APPLICATION NUMBER: 60/266,767

PRIOR APPLICATION NUMBER: 60/267,057

PRIOR PILING DATE: 2001-02-07

REMAINING PATO: 2001-02-08

REMAINING PATO: 2001-02-08

REMAINING PATO: 2001-02-08

REMAINING PATO: 2001-02-08

REMAINING PATO: 2001-02-07

PRIOR PILING DATE: 2001-02-08

REMAINING PATO: 2001-02-08

REMAINING PATO: 2001-02-07

PRIOR APPLICATION NUMBER: 60/267,459

PRIOR FILING DATE: 2001-02-08

REMAINING PATO: 2001-02-08

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REMAINING PATO: 200
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Rastelli, Luca
Miller, Charles E.
Gerlach, Valerie
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RIHLDWKVLFCHVHLVSIFLSALNSSANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPE
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Grosse, William M.
Alsobrook II, John P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Colman, Steven D. Wolenc, Adam R. Pena, Carol E. A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lepley, Denise M.
Rieger, Daniel K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gusev, Vladimir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shimkets, Richard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Zerhusen, Bryan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Application US/10072012
b. US20040033493A1
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Vladimir Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Velizar
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                                                                                                                               File Wrapper or
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APPLICANT: Alsobrook II, John P.
APPLICANT: Lepley, Denise M.
APPLICANT: Rieger, Daniel K.
APPLICANT: Burgess, Catherine E.
TITLE OF INVENTION: Proteins and Nucleic Acids Enc.
FILE REFERENCE: 21402-258
CURRENT APPLICATION NUMBER: US/10/072,012
CURRENT FILING DATE: 2002-01-31
PRIOR APPLICATION NUMBER: 60/265,102
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: 60/265,514
PRIOR APPLICATION NUMBER: 60/265,517
PRIOR APPLICATION NUMBER: 60/265,517
PRIOR APPLICATION NUMBER: 60/265,517
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PRIOR APPLICATION NUMBER: 60/265,517
PRIOR APPLICATION NUMBER: 60/265,517
PRIOR APPLICATION NUMBER: 60/265,517
PRIOR APPLICATION NUMBER: 60/265,512
PRIOR FILING DATE: 2001-01-31
PRIOR APPLICATION NUMBER: 60/265,412
PRIOR APPLICATION NUMBER: 60/265,395
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US-10-072-012-534
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Best Local S
Matches 313
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Taupier Jr, Raymond J
Gusev, Vladimir Y.
Colman, Steven D.
Wolenc, Adam R.
Pena, Carol B. A
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Spytek, Kimberly
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Shimkets, Richard
                                                                                                                                                                                                                                                                                                                                                                                                                 Furtak, Katarzyna
Grosse, William M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rastelli, Luca
Miller, Charles E.
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Anderson, David W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gangolli, Esha
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Pred. No. 2.5e-139;
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                                                                                                                                                                                                                                                                                              Encoding
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Taupier

APPLICANT: APPLICANT: APPLICANT:

Padigaru,

APPLICANT:

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APPLICANT: Zylka, Mark
APPLICANT: Simon, Melvin
APPLICANT: Simon, Melvin
APPLICANT: Han, Sang-kyou
FIITLE OF INVENTION: PAIN SIGNALING MOLECULES
FILE REFERENCE: CALTE.4C1CP1C1
CURRENT APPLICATION NUMBER: US/10/957,135
CURRENT FILING DATE: 2004-09-30
PRIOR APPLICATION NUMBER: US 60/222,344
PRIOR FILING DATE: 2000-08-01
PRIOR FILING DATE: 2000-05-04
PRIOR PILING DATE: 2000-05-04
PRIOR APPLICATION NUMBER: US 09/704,707
PRIOR PILING DATE: 2000-11-03
PRIOR PILING DATE: 2001-04-19
PRIOR APPLICATION NUMBER: US 60/285,493
PRIOR APPLICATION NUMBER: US 09/849,869
PRIOR FILING DATE: 2001-05-04
PRIOR FILING DATE: 2001-05-04
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US-10-957-135-31
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; ORGANISM: Homo sapiens
US-10-072-012-534
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Best Local Similarity 97.2
                                                                                                                                                                                                                                                                                                                                                                          Sequence 31, Application US/10957135
Publication No. US20050037468A1
GENERAL INFORMATION:
APPLICANT: Anderson, David J.
APPLICANT: Dong, Xinzhong
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SEQ ID NO 534
LENGTH: 322
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PRIOR APPLICATION NUMBER: 60/266,767
PRIOR APPLICATION NUMBER: 60/266,767
PRIOR APPLICATION NUMBER: 60/267,057
PRIOR PILING DATE: 2001-02-07
PRIOR PILING DATE: 2001-02-07
PRIOR PILING DATE: 2001-02-07
PRIOR PILING DATE: 2001-02-07
PRIOR PILING DATE: 2001-02-07
PRIOR PILING DATE: 2001-02-08
PRIOR PILING DATE: 2001-02-08
PRIOR PILING DATE: 2001-02-08
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PRIOR PILING DATE: 2001-02-08
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Pred. No. 2.5e-139;
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PRIOR APPLICATION NUMBER: US 10/183,116
PRIOR FILING DATE: 2002-06-26
NUMBER OF SEQ ID NOS: 109
SOFTWARE: FRSTSEQ for Windows Version 4.
SEQ ID NO 31
LENGTH: 322
TYPE: PRT
ORGANISM: Homo sapiens
US-10-957-135-31
       FILE OF INVESTION: FAIR SECURIOR SPILE REFERENCE: CALTE OOG 1

CURRENT FILING DATE: 2005-03-17

PRIOR APPLICATION NUMBER: US 09/849,869

PRIOR PILING DATE: 2001-05-04

PRIOR PILING DATE: 2001-05-04

PRIOR APPLICATION NUMBER: US 60/222,344

PRIOR FILING DATE: 2000-08-01

PRIOR APPLICATION NUMBER: US 60/202,027

PRIOR APPLICATION NUMBER: US 60/202,027

PRIOR PILING DATE: 2000-05-04

PRIOR PILING DATE: 2000-11-03

PRIOR APPLICATION NUMBER: US 09/704,707

PRIOR PILING DATE: 2000-11-03

PRIOR PILING DATE: 2000-14-03

PRIOR APPLICATION NUMBER: US 60/285,493

PRIOR FILING DATE: 2001-04-19

NUMBER OF SEQ ID NOS: 115

SOFTWARE: FASTSEQ for Windows Version 4.0

SEQ ID NO 31

LENGTH: 322

TYPE: PRT

ORGANISM: Homo sapiens
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US-11-083-611-31
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Best Local S
Matches 313
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                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Anderson, David J.
APPLICANT: Dong, Xinzhong
APPLICANT: Zylka, Mark
APPLICANT: Simon, Melvin
                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Han, Sang-kyou
TITLE OF INVENTION: PAIN SIGNALING MOLECULES
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97.2%;
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Pred. No. 2.5e-139;
5; Mismatches 4;
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Query Match Best Local Similarity

97.1%; 97.2%;

Score 1642; DB 6; Pred. No. 2.5e-139;

Length 322;

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Sequence 2, Application US/09867570

Publication No. US20040076951A1

GENERAL INFORMATION:
APPLICANT: WEI, Ming-Hui et al.
TITLE OF INVENTION: ISOLATED HUMAN G-PROTEIN COUPLED
TITLE OF INVENTION: RECEPTORS, NUCLEIC ACID MOLECULES ENCY
TITLE OF INVENTION: RECEPTORS, NUCLEIC ACID MOLECULES ENCY
TITLE OF INVENTION: RECEPTORS, AND USES THEREOF
FILE REFERENCE: CL000900-CLP
CURRENT APPLICATION NUMBER: US/09/867,570

CURRENT APPLICATION NUMBER: US/99/867,570

CURRENT APPLICATION NUMBER: 09/695,045
PRIOR APPLICATION NUMBER: 09/695,045
PRIOR FILING DATE: 2000-10-25
NUMBER OF SEQ ID NOS: 4
SOPTWARE: FRASEEQ FOR Windows Version 4.0

SEQ ID NO 2
LENGTH: 337
TYPE: PRT
CORGANISM: Human
US-09-867-570-2
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                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
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Pred. No. 2.6e-139;
5; Mismatches 4;
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Sequence 2, Application US/10401397A
Publication No. US20030212001A1
GENERAL INFORMATION:
APPLICANT: Peri, Krishna G.
APPLICANT: Moffett, Serge
APPLICANT: Abran, Daniel
TITLE OF INVENTION: METHODS AND COMPOUNDS FOR PRE
TITLE OF INVENTION: INTRAOCULAR PRESSURE AND REI
FILE REFERENCE: 4518/1M674US1
CURRENT APPLICATION NUMBER: US/10/401,397A
CURRENT FILING DATE: 2003-03-27
PRIOR APPLICATION NUMBER: US 60/367,513
PRIOR APPLICATION NUMBER: US 60/367,513
PRIOR APPLICATION NUMBER: US 60/367,513
PRIOR FILING DATE: 2002-03-27
NUMBER OF SEQ ID NOS: 8
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TITLE OF INVENTION: Determination of a ligand
FILE REFERENCE: P03-0006PCT
CURRENT APPLICATION NUMBER: US/10/505,486
CURRENT FILING DATE: 2004-08-20
PRIOR APPLICATION NUMBER: UP 2002-45728
PRIOR FILING DATE: 2002-02-22
PRIOR APPLICATION NUMBER: UP 2002-213949
PRIOR FILING DATE: 2002-07-23
PRIOR APPLICATION NUMBER: UP 2002-213949
PRIOR FILING DATE: 2002-07-13
PRIOR APPLICATION NUMBER: UP 2002-298237
PRIOR FILING DATE: 2002-10-11
NUMBER OF SEQ ID NOS: 233
SEQ ID NO 104
LENGTH: 560
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US-10-401-397A-2
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US-10-505-486-104
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Best Local S
Matches 313
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97.2%;
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Pred. No. 4.5e-139;
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APPLICANT: Peri, Krishna G.

APPLICANT: Moffett, Serge
APPLICANT: Moffett, Serge
APPLICANT: Abran, Daniel
ITILE OF INVENTION: METHODS AND COMPOUNDS FOR PREVENTION AND TREATMENT OF ELEVATED
ITILE OF INVENTION: INTRACCULAR PRESSURE AND RELATED CONDITIONS
FILE REFERENCE: 04518/100M674-US2
CURRENT APPLICATION NUMBER: US/10/977,810
CURRENT FILING DATE: 2004-10-28
PRIOR APPLICATION NUMBER: US 60/367,513
PRIOR FILING DATE: 2002-03-27
NUMBER OF SEQ ID NOS: 8
SOFTWARE: PatentIn version 3.1
SEQ ID NO 2
LENGTH: 322
TYPE: PRT
ORGANISM: Homo sapiens
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US-10-977-810-2
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US-10-401-397A-2
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SEQ ID NO 2
LENGTH: 322
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Best Local Similarity 96.9
Matches 312; Conservative
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CLSILWPIWYHCRRPRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWCETSDFIT
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96.9%; Pred. No. 7e-139;
tive 6; Mismatches 4; Indels
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Pred. No. 7e-139;
6; Mismatches 4; Indels
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VDEGGGWLPQETLELSGSRLEQ 322
              VDEGGGWLPQETLELSGSKLEQ 322
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Search completed: February 3, 2006, 20:46:57
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Result
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/cgn2_6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*
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APPLICATION NUMBER: 60/184, FILLING DATE: 2000-02-23 APPLICATION NUMBER: 60/188, FILLING DATE: 2000-03-13 APPLICATION NUMBER: 60/217, FILLING DATE: 2000-07-11 APPLICATION NUMBER: 60/217, FILLING DATE: 2000-07-11 APPLICATION NUMBER: 60/217, FILLING DATE: 2000-07-20 APPLICATION NUMBER: 60/218, FILLING DATE: 2000-07-20 APPLICATION NUMBER: 60/218, FILLING DATE: 2000-07-20 APPLICATION NUMBER: 60/218, FILLING DATE: 2000-07-20 APPLICATION NUMBER: 60/218, FILLING DATE: 2000-07-20 APPLICATION NO 111 APPLICATION NO 111 APPLICATION NO 111 APPLICATION SAPIENS HOMO SAPIE	STANDARD STANDARD CONTRACTOR CONT	111.3 111.2 110.4 110.4 110.2 10.0 10.0 10.0 10.0 10.0 10.0 10.
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Query Match Best Local Similarity

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Score 485.5; DB 6; Pred. No. 3.5e-35;

Length 311;

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NUMBER OF SEQ ID NOS: 184
SOFTWARE: PatentIn version 3.0
SEQ ID NO 62
             Query Match
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Matches
                                                                      ORGANISM: Homo sapiens -10-980-388-62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Novel G Protein-Coupled Receptors Cross-Reference To FILE REFERENCE: 00325.US1
CURRENT APPLICATION NUMBER: US/10/980,388
CURRENT FILING DATE: 2004-11-02
PRIOR APPLICATION NUMBER: US/09/791,932
PRIOR PILING DATE: 2001-02-23
PRIOR APPLICATION NUMBER: 60/184,305
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OR APPLICATION NUMBER: 60/184,397

OR FILING DATE: 2000-02-23

OR APPLICATION NUMBER: 60/184,247

OR FILING DATE: 2000-02-23

OR APPLICATION NUMBER: 60/188,880

OR APPLICATION NUMBER: 60/188,880
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APPLICATION NUMBER: 60/184,304
FILING DATE: 2000-02-23
APPLICATION NUMBER: 60/184,303
FILING DATE: 2000-02-23
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Huff, Rita M.
Wood, Linda S.
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Hiebsch, Ronald R.
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28.7%; Score 485.5; DB 6; 38.5%; Pred. No. 5.8e-35; tive 51; Mismatches 120;
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                                                                         ; ORGANISM: Homo US-10-980-388-113
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                 Query Match
Best Local Similarity
                                                                                                                                                 SOFTWARE: PatentIn version SEQ ID NO 113
   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Novel G Protein-Coupled Receptors Cross-Reference FILE REFERENCE: 00325.US1
CURRENT APPLICATION NUMBER: US/10/980,388
CURRENT FILING DATE: 2004-11-02
                                                                                                                                                                                    Remaining Prior Application data removed - See File Wrapper or PALM NUMBER OF SEQ ID NOS: 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US/09/791,932
PRIOR FILING DATE: 2001-02-23
PRIOR APPLICATION NUMBER: 60/184,305
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PRIOR FILING DATE: 2000-07-20
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                                                                                                            LENGTH: 31
TYPE: PRT
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APPLICATION NUMBER: 60/184,247
FILING DATE: 2000-02-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 2000-02-23
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Huff, Rita M.
Wood, Linda S.
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   Conservative
28.4%; Score 479.5; DB 6; 38.2%; Pred. No. 1.1e-34; tive 51; Mismatches 121;
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RESULT 4
US-10-055-877-244
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TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids:
FILE REFERENCE: 21402-251
CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR PILING DATE: 2001-01-23
PRIOR PILING DATE: 2001-01-23
PRIOR FILING DATE: 2001-01-23
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OR APPLICATION NUMBER: 60/264,117
OR FILING DATE: 2001-01-25
OR APPLICATION NUMBER: 60/264,139
OR FILING DATE: 2001-01-25
OR APPLICATION NUMBER: 60/264,478
OR FILING DATE: 2001-01-26
OR APPLICATION NUMBER: 60/263,351
OR APPLICATION NUMBER: 60/272,870
OR APPLICATION NUMBER: 60/272,870
OR FILING DATE: 2001-03-02
                                                                                                                                                                APPLICATION NUMBER: 60/263,799 FILING DATE: 2001-01-24
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Taupier Jr., Raymond
Pena, Carol
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Shimkets, Richard
Gusev, Vladimir
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Miller, Charles
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Burgess, Cahterine
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Spytek, Kimberly
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Kekuda, Ramesh
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ısen, Bryan
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RESULT 5
US-10-055-877-245
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; ORGANISM: Homo sapiens
US-10-055-877-244
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SEQ ID NO 244
LENGTH: 343
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Best Local Similarity
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PRIOR FILING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/275,927
PRIOR FILING DATE: 2001-03-14
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Invention: Novel Polypeptides and Nucleic Acids Encoded Thereby
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                                                                                      Vernet, Corine
Taupier Jr., Raymond
Pena, Carol
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Shimkets, Richard
Gusev, Vladimir
                                   Casman,
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Mezes, Peter
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Kekuda, Ramesh
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Burgess, Cahterine
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                                    Stacie
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; SOFTWARE: PatentIn Ver. 2
; SEQ ID NO 245
; LENGTH: 343
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-055-877-245
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CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR FILING DATE: 2001-01-19
PRIOR FILING DATE: 2001-01-29
PRIOR PPLICATION NUMBER: 60/263,598
PRIOR APPLICATION NUMBER: 60/263,799
PRIOR FILING DATE: 2001-01-24
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR APPLICATION NUMBER: 60/264,139
PRIOR APPLICATION NUMBER: 60/264,139
PRIOR FILING DATE: 2001-01-25
PRIOR FILING DATE: 2001-01-25
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PRIOR FILING DATE: 2001-01-30
PRIOR PEPLICATION NUMBER: 60/272,870
PRIOR WILING DATE: 2001-03-02
PRIOR APPLICATION NUMBER: 60/275,990
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
                                                                                                                                                                                                                     Sequence 71, Application US/10055877
Publication No. US20050288241A1
GENERAL INFORMATION:
APPLICANT: DeCristofaro, Marc
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Best Local Similarity 34.6%; Pred. No. 1.2e-28;
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PRIOR FILING DATE: 2001-01-26
                                           APPLICANT:
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Gerlach, Valerie
Spytek, Kimberly
Ratelli, Luca
Kekuda, Ramesh
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                                                                                                                                                     Padigaru, Muralidhara
Miller, Charles
Tchernev, Velizar
                                                                 Anderson, David
Ballinger, Robert
Gerlach, Valerie
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PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR FILING DATE: 2001-01-23
PRIOR APPLICATION NUMBER: 60/263,799
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR FILING DATE: 2001-01-25
PRIOR FILING DATE: 2001-01-25
PRIOR FILING DATE: 2001-01-25
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SEQ ID NO 71
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Best Local Similarity
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TITLE OF INVENTION: Novel Polypeptides and
FILE REFERENCE: 21402-51
CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 202-01-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Remaining Prior Application data removed - See File Wrapper or PALM NUMBER OF SEQ ID NOS: 512
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 343
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SANPIIYFFVGSFRQRQNRQNLKLVLQRALQDTPEVDEGGGWLPQE-TLEL
                                                                               KMPLT-RLYVTILLTVLVFLLCGLPFGIQWALFSRIHLDW--KVLFCHVHLVSIFLSALN
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Shimkets, Richard
Gusev, Vladimir
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Taupier Jr., Raymond
Pena, Carol
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Patturajan, Meera
Burgess, Cahterine
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Eisen, Andrew
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Andrew, David
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usen, Bryan
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34.2%; Pred. No. 4.7e-28;
ative 53; Mismatches 118
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NUMBER OF SEQ ID NOS: 184
SOFTWARE: PatentIn version 3.0
SEQ ID NO 119
LENGTH: 323
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PRIOR APPLICATION NUMBER: 60/217,369
PRIOR FILING DATE: 2000-07-11
PRIOR APPLICATION NUMBER: 60/217,370
PRIOR FILING DATE: 2000-07-11
PRIOR APPLICATION NUMBER: 60/218,492
PRIOR FILING DATE: 2000-07-20
PRIOR FILING DATE: 2000-07-20
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CURRENT APPLICATION NUMBER: US/10/980,388
CURRENT FILING DATE: 2004-11-02
PRIOR APPLICATION NUMBER: US/09/791,932
PRIOR FILING DATE: 2001-02-23
PRIOR FILING DATE: 2001-02-23
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PRIOR FILING DATE: 2000-02-23
PRIOR APPLICATION NUMBER: 60/184,304
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PRIOR FILING DATE: 2000-02-23
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ORGANISM: Homo sapiens
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APPLICATION NUMBER: 60/188,880
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APPLICATION NUMBER: 60/184,247
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    278
                                                                                                                                                                                  169 PAVPLPANACGLLRNSACPLVCPRYHVASVTWFLVLARVAWTAGVVLFVWVTCCSTR-PR
                                                                                                                                                                                                                           152 LRSILEWMFCDFLFSGANSVWCETSDFITIAWLVFLCVVLCGSSLVLLVRILCGSRKMPL 211
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                                                                                                                                                                                                                                                                                                                                                                     52
                                                                                                                                                                                                                                                                                                                   93 PISKIĻSPYMTEPYE-IGLSMĻSAISTERCLSILWPIWYHCRRPRYLSSVMCVLLWALSL 151
                                                                                                                                                                                                                                                                        GAQDTLYEVLTELMEAVGLMLLAAFSVERCLSDLFPACYQGCRERHASAVLCALVMTPTL 168
SSKPLIYSGLG--ROPGKREPLRSVLRRALGEGAELGARGOSLPMGLL 323
                                        SANPITYFFVGSFRQRQNRQNLKLVLQRALQDTPEVDEGGGWLPQETL 313
                                                                                          PRLYGIVLGALLLLFFCGLPSVFYWSLQPLLNF-
                                                                                                                                   TRLYVTILLTVLVFLLCGLPFGIQWALFSRIHLDWKVLFCHVHLVSIF-----LSALNS 265
                                                                                                                                                                                                                                                                                                                                                                 LTLIVGLGGPVGNGLVLWNLGFRIKKGPFSIYLLHLAAADFLFLSCRVGFS---VAQAAL 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ruff, Valerie
Huff, Rita M.
Wood, Linda S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Score 375.5; DB 6; Length 323; pred. No. 1.2e-25; 48; Mismatches 115; Indels 23
                                                                                        LLPVFSPLATLLACVNS
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-055-877-246
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US-10-055-877-246
Query Match
Best Local Similarity
Matches 96; Conserv
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                                                                                                                          SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 246
LENGTH: 342
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CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
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PRIOR APPLICATION NUMBER: 60/263,598
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PRIOR APPLICATION NUMBER: 60/264,117
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PRIOR FILING DATE: 2001-01-24
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                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 60/272,870 FILING DATE: 2001-03-02
                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 60/264,478
                                                                                                                                                                                                                 FILING DATE: 2001-03-1
                                                                                                                                                                                                                                                                                                                      FILING DATE:
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Andrew, Peter
Mezes, Peter
Patturajan, Meera
Patturajan, Cahterine
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Miller, Charles
Tchernev, Velizar
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Taupier Jr., Raymond
Pena, Carol
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Shimkets, Richard
Gusev, Vladimir
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Wolenc, Adam
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Kekuda, Ramesh
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Spytek, Kimberly
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Gerlach, Valerie
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     21.7%; Score 367.5; DB 6; ilarity 32.4%; Pred. No. 6.1e-25; Conservative 61; Mismatches 110;
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       Indels 29;
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         FILE REFERENCE: 21402-51
CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR FILING DATE: 2001-01-23
PRIOR PILING DATE: 2001-01-23
PRIOR FILING DATE: 2001-01-24
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PRIOR APPLICATION NUMBER: 60/264,139
PRIOR FILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,139
PRIOR PILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,478
PRIOR APPLICATION NUMBER: 60/264,478
PRIOR APPLICATION NUMBER: 60/263,351
PRIOR FILING DATE: 2001-01-26
PRIOR FILING DATE: 2001-01-26
PRIOR APPLICATION NUMBER: 60/263,351
PRIOR FILING DATE: 2001-01-30
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TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
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Miller, Charles
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Shimkets, Richard
Gusev, Vladimir
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Mezes, Peter
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Kekuda, Ramesh
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Spytek, Kimberly
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Andrew, David
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o. US20050288241A1
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NUMBER: 60/272,870
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; LENGTH: 319
; TYPE: PRT
; ORGANISM: Mus musculus
US-10-055-877-247
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PRIOR APPLICATION NUMBER: 60/275,990
PRIOR PILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 512
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 247
                                                                                                                                                                                                                                                                                                                                                                   Matches
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Best Local Similarity
                                                                                                                                              142
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204
                                261 SALNSSANPITYFFVGSFRORONRONLKLVLORALODTPEVDEGGGWLPQE-TLEL 315
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                                                                                                                                                                                                                                                                                                                               36 IISLVALTGNAVVLWLLGCRMRRNAVSIYILNLVAANFLFLSGHIIFSPLPLIN-----
                                                                                                                                                                                                                                                                                             29 LLCLCGLVGNGLVLWFFGFSIKRTPFSIYFLHLASADGMYLFSKAV---IALLNMGTFLG
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                                                                                                      CGSRKMPLT-RLYVTILLTVLVFFLCGLPFGIQWALFSRIHLDW--KVLFCHVHLVSIFL
                                                                                                                                            LLWMLSFLVTSIHNYFCMFLGHEAPGTVCRNMDIALGILLFFLFCPLMVLPCLALILHVE
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                                                                                                                                                                                                                                                       ----IRHPISKILSPVMTFPYFIGLSMLSAISTERCLSILWPIWYHCRRPRYLSSVMCV 144
                                                                      CRARRORSAKLNHVVLAIVSVÉLVSSIYLGIDWELE-----WVFQIPAPEPEYVTDLC
                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                     21.7%; Score 366.5; DB 6; 33.1%; Pred. No. 6.9e-25;
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US-10-055-877-69 RESULT 10 GENERAL INFORMATION:
APPLICANT: DeCristofaro, Marc Sequence 69, Application US/10055877 Publication No. US20050288241A1 APPLICANT:
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OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby

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US-10-980-388-39

Sequence 39, Application US/10980388

Dublication No. US20050255490A1
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Best Local Similarity
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APPLICANT: Parod
APPLICANT: Hiebs
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PRIOR APPLICATION NUMBER: 60/275,927
PRIOR FILING DATE: 2001-03-14
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PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: 60/263,598
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PRIOR APPLICATION NUMBER: 60/264,139
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CURRENT FILING DATE: 2002-01-22
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REFERENCE: 00325.US1
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APPLICATION NUMBER: 60/263,799
FILING DATE: 2001-01-24
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                 INVENTION: Novel G Protein-Coupled Receptors Cross-Reference
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                                                                                                           Parodi, Luis A.
Hiebsch, Ronald R.
Lind, Peter
                                 Huff, Rita M. Wood, Linda S.
                                                                  Ruff, Valerie
                                                                                       Kaytes,
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                                                                                         Paul S.
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Pred. No. 2e-23;
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APPLICANT: Vogeli, Ge
APPLICANT: Parodi, I
APPLICANT: Hiebsch,
APPLICANT: Lind, Pet
APPLICANT: Kaytes, E
APPLICANT: Ruff, Val
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Best Local
            PRIOR APPLICATION NUMBER: US/09/791,932
PRIOR FILING DATE: 2001-02-23
PRIOR APPLICATION NUMBER: 60/184,305
PRIOR FILING DATE: 2000-02-23
PRIOR PPLICATION NUMBER: 60/184,304
PRIOR FILING DATE: 2000-02-23
PRIOR APPLICATION NUMBER: 60/184,303
                                                                                                                                                       TITLE OF INVENTION: Novel G Protein-Coupled Receptors Cross-Reference To Related Appli
FILE REFERENCE: 00325.US1
CURRENT APPLICATION NUMBER: US/10/980,388
CURRENT FILING DATE: 2004-11-02
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CURRENT FILING DATE: 2004-11-02
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PRIOR FILING DATE: 2001-02-23
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FILING DATE: 2000-02-23
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57; Conserv
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APPLICANT: Kaytes, Paul S.
APPLICANT: Ruff, Valerie
APPLICANT: Huff, Rita M.
APPLICANT: Wood, Linda S.
TITLE OF INVENTION: Novel G Protein-Coupled Receptors Cross-Reference To Related Appl CURRENT APPLICATION NUMBER: US/10/980,388
CURRENT FILING DATE: 2004-11-02
PRIOR APPLICATION NUMBER: US/09/791,932
PRIOR FILING DATE: 2001-02-23
PRIOR APPLICATION NUMBER: US/09/791,932
PRIOR APPLICATION NUMBER: US/09/791,932
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; LENGTH: 187
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-980-388-98
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PRIOR FILING DATE: 2000-02-23
PRIOR PELICATION NUMBER: 60/184,304
PRIOR PELICATION NUMBER: 60/184,304
PRIOR PELICATION NUMBER: 60/184,303
PRIOR PELICATION NUMBER: 60/184,397
PRIOR APPLICATION NUMBER: 60/184,397
PRIOR APPLICATION NUMBER: 60/184,397
PRIOR APPLICATION NUMBER: 60/184,247
PRIOR APPLICATION NUMBER: 60/184,247
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PRIOR FILING DATE: 2000-07-20
Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 184
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PRIOR APPLICATION NUMBER: 60/188,880
PRIOR FILING DATE: 2000-03-13
PRIOR APPLICATION NUMBER: 60/217,369
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OR APPLICATION NUMBER: 60/184,397
OR FILING DATE: 2000-02-23
OR APPLICATION NUMBER: 60/184,247
OR FILING DATE: 2000-02-23
OR APPLICATION NUMBER: 60/188,880
OR FILING DATE: 2000-03-13
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FILING DATE: 2000-02-23
APPLICATION NUMBER: 60/184,247
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APPLICATION NUMBER: 60/217,370
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                       APPLICATION NUMBER: 60/217,369
FILING DATE: 2000-07-11
APPLICATION NUMBER: 60/217,370
FILING DATE: 2000-07-11
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; LENGTH: 211
; TYPE: PRT
; ORGANISM: Homo 8
US-10-980-388-97
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CURRENT FILING DATE: 2004-12-21
PRIOR APPLICATION NUMBER: US 09/101,518
PRIOR FILING DATE: 1998-12-21
PRIOR APPLICATION NUMBER: PCT/US96/00499
PRIOR FILING DATE: 1996-01-11
NUMBER OF SEQ ID NOS: 9
SOFTWARE: PatentIn version 3.1
SBQ ID NO 9
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NUMBER OF SEQ ID NOS: 184
SOFTWARE: PatentIn version 3.0
SEQ ID NO 97
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Publication No. US20060014243A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 14.4%; Score 243; DB 6; Best Local Similarity 38.4%; Pred. No. 2.2e-14; Matches 61; Conservative 26; Mismatches 68
                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/218,492 PRIOR FILING DATE: 2000-07-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Li, Yi
TITLE OF INVENTION: Human G-Protein Chemokine
FILE REFERENCE: PF218C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 353
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               109 GAQDTLYFVLTFLWFAVGLWLLAAFSVERCLSDLFFACYQGCRPRHASAVLCALVWTPTL
                                                                        195 GNNTANWRMLLRILPQSFGFIVPLLIMLFCYGFTLRTLFKAHMGQKHRAMRVIFAVVLIF
                                                                                                                                                      140
                                                                                                                                                                                      122 LSILWPIWYHCRR----PRYLSSVMCVLLWALSLLRSILEWMFCDFLFSGANSVWC-ETS 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         52 LTLIVGLGGPVGNGLVLMNLGFRIKKGPFSIYLLHLAAADFLFLSCRVGFS---VAQAAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             33 LTCIISLVALTGNAVVLWLLGCRMRRNAVSIYILNLVAANFLFLSGHIIFSPLPLINIRH
                                                                                                                                                                                                                              89 LLFALTLPIWAASKVNGWIFGTFL------CKVVSLLKEVNFYSGILLLACISVDRY 139
                                                                                                                                                                                                                                                                                                        93
                                                                                                                                                                                                                                                                73 FLF-----LSGHIIFSPLPLINIRHPISKILSPVMTFPYFTGLSMLSAISTERC 121
                                                                                                                                                                                                                                                                                                                                               19 EETPCYNQTLSFTG-----LTCIISLVALTGNA-VVLWLLGCRMRRNAVSIYILNLVAAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             353
                                                                                                                                                                                                                                                                                                                                                                                       81;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PISKILSPYMTEPYE-IGLSMLSAISTERCLSILWPIWYHCRRPRYLSSYMCVLLWALSL 151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PAVPLPANACGLIRNSACPLVCPRYHVASVTWFLVLARV 207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LRSILEWMFCDFLFSGANSVWCETSDFITIAWLVFLCVV 190
                                                                                                              DFITIAMLVFLCVV--LCGSSLVLLVRILC-GSRKMPLTRLYV-----TILLTVLVF
                                                                                                                                                    LAIV-----HATRTLTQKRYLVKFICLSIWGLSLLLALPVLLFRRTVYSSNVSPACYEDM 194
                                      LLCGLPFG-----
LLCWLPYNLVLLADTLMRTQVIQETCERRNHIDRALDATEI-----LGILHSCLNPLIY
                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sapiens
                                                                                                                                                                                                                                                                                                                                                                                     12.7%; Score 215.5; DB 7; 26.3%; Pred. No. 8.7e-12; tive 55; Mismatches 105;
                                  IQWALFSRIHLDWKVLFCHVHLVSIFLSALNSSANPIIY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         68; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Receptor HSATU68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 211
                                                                                                                                                                                                                                                                                                                                                                                                                               Length 353;
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                                                                                                                                                                                                                                                                                                                                                                                             67;
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                                            272
        308
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273 FFVG-SFR 279 |:| || 309 AFIGQKFR 316

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FILE REPERENCE: 21402-251
CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR FILING DATE: 2001-01-19
PRIOR FILING DATE: 2001-01-23
PRIOR PRIOR PRIOR DATE: 2001-01-24
PRIOR FILING DATE: 2001-01-24
PRIOR FILING DATE: 2001-01-24
PRIOR FILING DATE: 2001-01-25
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US-10-055-877-225
                                       Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 512
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 225
LENGTH: 259
TYPE: PAT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Publication No.
                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/272,870
PRIOR FILING DATE: 2001-03-02
PRIOR APPLICATION NUMBER: 60/275,990
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2001-01-07
PRIOR APPLICATION NUMBER: 60/263,351
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 60/272,870
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APPLICANT:
APPLICANT:
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TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
FILE REFERENCE: 21402-251
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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APPLICANT: Padigaru, Muralidhara
APPLICANT: Miller, Charles
APPLICANT: Tchernev, Velizar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
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OTHER INFORMATION: Description of Artificial Sequence: 7tm_1 domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PPLICANT:
                           EATURE:
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                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 60/264,478 FILING DATE: 2001-01-26
                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 2001-01-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mezes, Peter
Patturajan, Meera
Burgess, Cahterine
Eisen, Andrew
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Taupier Jr., Raymond
Pena, Carol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shenoy, Suresh
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Shimkets, Richard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Andrew,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Guo, Xiaojia
Zerhusen, Bryan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ratelli, Luca
Kekuda, Ramesh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gerlach, Valerie
Spytek, Kimberly
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Zhong, Mei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Application US/10055877
D. US20050288241A1
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                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 29.2
""""
Conservative
226 IMSSTCELERVLPTALLVTLWLAYVNSCLNPIIY
                                   250
                                                               172 GPLLPLLVILVCYTRILRTLRKAAKTLLVVVVVVFVLCWLPY-----FIVLLLDTLCLSI 225
                                                                                                                               117 LLSL-----PPLLFSWVKTVEEGNGTLNVNVTVCLIDFPEESTASVSTWLRSYVLLSTLV 171
                                                                                              193 GSSLVILLVRILCGSRKM-PLTRLYVTILLTVLVFLLCGLPFGIQWALFSRIHLDWKVL--
                                                                                                                                                             152 LRSILEWMFCDFLFSGANSV--------WCETSDFITIAWL--VFLCVVLC 192
                                                                                                                                                                                             58 LCKLVTALDVVNMYASILLLTAISIDRYLAIVHPLRYRRRRTSPR-RAKVVILLVMVLAL 116
                                                                                                                                                                                                                             94 ISKILSPVMTFPYFIGLSMLSAISTERCLSILWPIWYHCRR--PRYLSSVMCVLLWALSL
                                                                                                                                                                                                                                                                                             44 GN-AVVLWLLGCRMRRNAVSIYILNLVAANFLFLSGHIIFSPLPLINI-----RHP
                                                                                                                                                                                                                                                             ----FCHVH-----LVSIFLSALNSSANPITY 272
                                                                                                                                                                                                                                                                                                                                            12.1%;
                                                                                                                                                                                                                                                                                                                             46;
                                                                                                                                                                                                                                                                                                                           Score 205; DB 6;
Pred. No. 5.2e-11;
6; Mismatches 88
 259
                                                                                                                                                                                                                                                                                                                                                         Length 259;
                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                             60;
                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                               249
                                                                                                                                                                                                                           151
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Search completed: February 3, 2006, 20:47:18
Job time : 17 secs

**Ipis** bade Blank (nsbto)

## November 2005

Published\_Applications Nucleic Acid and Published\_Applications Amino Acid database searches now generate two sets of results each. The Published\_Applications databases have been split into two parts to reduce the amount of time required for their daily updates. This results in more machine time being available for processing searches.

Newly published applications will appear in the Published\_Applications\_New databases; older published applications make up the Published\_Applications\_Main databases.

Searches run against Nucleic Acid Published\_Applications produce two sets of results, with the extensions '.rnpbm (Published\_Applications\_NA\_New).

Searches run against Amino Acid Published\_Applications produce two sets of results, with the extensions .rapbm (Published\_Applications\_AA\_New).

**IVI**? Lage Blank (uspto)